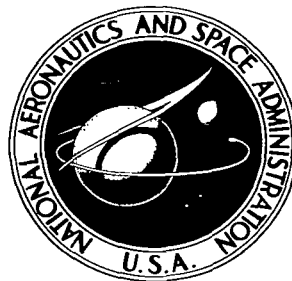


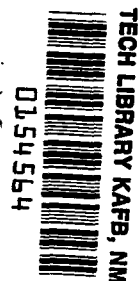
NASA TECHNICAL NOTE



NASA TN D-2085

C.1

LOAN COPY: RETL
AFWL (WLL-
KIRTLAND AFB, TX



NASA TN D-2085

FORTRAN PROGRAM FOR COMPUTING THE PRINCIPAL MOMENTS OF INERTIA OF A RIGID MOLECULE

by Janet G. Ehlers and Glenn R. Cowgill

*Lewis Research Center
Cleveland, Ohio*



**FORTTRAN PROGRAM FOR COMPUTING THE PRINCIPAL MOMENTS
OF INERTIA OF A RIGID MOLECULE**

By Janet G. Ehlers and Glenn R. Cowgill

**Lewis Research Center
Cleveland, Ohio**

NATIONAL AERONAUTICS AND SPACE ADMINISTRATION

**For sale by the Office of Technical Services, Department of Commerce,
Washington, D.C. 20230 -- Price \$1.25**

FORTRAN PROGRAM FOR COMPUTING THE PRINCIPAL MOMENTS
OF INERTIA OF A RIGID MOLECULE

By Janet G. Ehlers and Glenn R. Cowgill
Lewis Research Center

SUMMARY

A program to compute the principal moments of inertia of a rigid molecule was written in the FORTRAN language. This program eliminates considerations of symmetry and avoids hand calculations by reducing the input to structural parameters, that is, bond lengths and angles. The assignment of the structural parameters follows geometric models. In its present form the program is written for molecules with not more than 24 atoms.

INTRODUCTION

Many structurally complicated molecular species are becoming of interest at shock-tube and rocket-combustion temperatures. These include polyatomic species, such as HBO_2 and BFCl , which have no axes of symmetry, and polymers, such as $(\text{NaOH})_2$ and $(\text{HBO}_2)_3$. Obtaining thermodynamic functions for these species requires that the principal moments of inertia or the product of the principal moments be known. The calculation, however, becomes cumbersome for large molecules or for molecules in which the center of mass and the directions of the principal axes through the center of mass are not immediately apparent. In order to eliminate considerations of symmetry and to avoid hand calculations, the method of reference 1 together with the authors' technique to minimize the input was programmed in the FORTRAN language.

The program described herein was used to compute the moments of inertia reported in reference 2. In its present form the program is written for molecules with not more than 24 atoms; however, extension to consider molecules with more than 24 atoms could be easily accomplished. Any set of atomic weights with the corresponding value of Avogadro's number may be selected as input to the program.

The program decks are available from the authors upon request in either FORTRAN II or FORTRAN IV. Also included for the user's convenience are data cards for the atomic weights based on the chemical scale of natural oxygen ($\text{O} = 16.0000$) and the 1961 Table of Atomic Weights (see ref. 3) based on the exact weight of 12 for carbon 12.

SYMBOLS

A	symmetric matrix
I_{xx}, I_{yy}, I_{zz}	moments of inertia
$I_{xy}, I_{yz}, I_{xz},$ I_{yx}, I_{zy}, I_{zx}	products of inertia
M	$\sum_i m_i$
m_i	mass of the i^{th} atom
r	bond length
$r_{i,k}$	bond length between the i^{th} and k^{th} atoms
X,Y	atoms in pyramidal or tetrahedral molecules
x_i, y_i, z_i	Cartesian coordinates of the i^{th} atom
α	angle
θ_i	branch angle with vertex at the i^{th} atom
λ	eigenvalue
σ_i	out-of-plane-atom angle with vertex at the i^{th} atom
ϕ_i, ϕ_k	chain angles with vertex at the i^{th} and k^{th} atoms

METHOD OF CALCULATION

The principal moments of inertia are the eigenvalues of the real symmetric matrix A

$$A = \begin{pmatrix} I_{xx} & I_{xy} & I_{xz} \\ I_{yx} & I_{yy} & I_{yz} \\ I_{zx} & I_{zy} & I_{zz} \end{pmatrix}$$

The diagonal elements are the moments of inertia about the center of mass; the off-diagonal elements are the products of inertia about the center of mass.

The matrix elements are calculated according to the equations given in reference 1. These equations, where m_i is the mass of the i^{th} atom, x_i, y_i, z_i

are the Cartesian coordinates of this atom, and $M = \sum_i m_i$, are reproduced for convenience:

$$I_{xx} = \sum_i m_i (y_i^2 + z_i^2) - \frac{1}{M} \left(\sum_i m_i y_i \right)^2 - \frac{1}{M} \left(\sum_i m_i z_i \right)^2$$

$$I_{yy} = \sum_i m_i (x_i^2 + z_i^2) - \frac{1}{M} \left(\sum_i m_i x_i \right)^2 - \frac{1}{M} \left(\sum_i m_i z_i \right)^2$$

$$I_{zz} = \sum_i m_i (x_i^2 + y_i^2) - \frac{1}{M} \left(\sum_i m_i x_i \right)^2 - \frac{1}{M} \left(\sum_i m_i y_i \right)^2$$

$$I_{xy} = I_{yx} = - \sum_i m_i x_i y_i + \frac{1}{M} \left(\sum_i m_i x_i \right) \left(\sum_i m_i y_i \right)$$

$$I_{xz} = I_{zx} = - \sum_i m_i x_i z_i + \frac{1}{M} \left(\sum_i m_i x_i \right) \left(\sum_i m_i z_i \right)$$

$$I_{yz} = I_{zy} = - \sum_i m_i y_i z_i + \frac{1}{M} \left(\sum_i m_i y_i \right) \left(\sum_i m_i z_i \right)$$

Any convenient set of orthogonal axes may be chosen to describe a molecule. The equations for the matrix elements give the moments of inertia and the products of inertia about a set of axes parallel to the assigned axes and with origin at the center of mass.

A convenient value in the calculation of thermodynamic properties is the product of the principal moments of inertia. This value is simply the determinant of the matrix A .

The three principal moments of inertia can be found individually as the roots λ of the secular equation

$$\begin{vmatrix} I_{xx} - \lambda & I_{xy} & I_{xz} \\ I_{yx} & I_{yy} - \lambda & I_{yz} \\ I_{zx} & I_{zy} & I_{zz} - \lambda \end{vmatrix} = 0$$

The derivation of this equation is discussed in texts on classical mechanics, such as reference 4.

An attempt to solve directly for the roots of the preceding cubic equation resulted in the appearance of false imaginary solutions, which were caused by inherent machine inaccuracies and rounding errors. Of several other possible methods, the iterative Jacobi method described in reference 5 was chosen to determine the principal moments inasmuch as this method does not depend on the existence of three unequal real roots. In the solution is found an orthogonal matrix, that is, an eigenvector matrix, which transforms the original symmetric matrix A into a diagonal matrix. This is accomplished through the annihilation of selected off-diagonal elements by elementary orthogonal transformations. The elements of the resulting diagonal matrix are the principal moments of inertia.

Because of rounding errors in the computation, the principal moments of inertia are accurate to approximately six significant figures. This degree of accuracy was determined by repeating computations for a molecule after a change in its orientation relative to the assigned axes. This accuracy, however, is adequate in the light of the reliability of experimental structural parameter data.

DISCUSSION OF THE PROGRAM FOR VARIOUS TYPES OF MOLECULES

Figure 1 is a flow diagram of the main program. The block numbers correspond to the statement numbers of the code listing. A flow diagram of the subroutine Jacobi is given in reference 5. The computation of the principal moments of inertia is simplified by equations for the Cartesian coordinates of atoms in terms of the appropriate bond lengths and angles. A molecule is classified according to its geometric type: planar, pyramidal, tetrahedral, other nonplanar, or general. The input for each geometric type follows a specified model. Although this input is reduced to bond lengths and angles, their order of assignment is thus not completely arbitrary.

Each of the six geometric types is illustrated in figure 2 with the equations for the atom coordinates given in table I. Sample computations for each type of input are given in table II. These computations can be used to check the program deck.

Table III is a listing of the code preceded by an explanation of the input cards, and followed by a listing of the input cards used for the computations in table II. The atomic weights and the corresponding Avogadro's number are based on the chemical scale of natural oxygen.

Type 1 - Planar

The input for planar molecules can be reduced to bond lengths and angles when the molecule is considered to be a chain of atoms with other atoms branching from the chain. Planar molecules without and with branching are illustrated in figures 2(a-1) and (a-2), respectively. Figure 2(a-1) represents a molecule such as HBO_2 , which has a chain of four atoms; figure 2(a-2) represents a molecule such as $(\text{HBO}_2)_3$, which can be visualized as a chain of six atoms with three two-atom branches attached to the chain. The branch atoms are specified according to the chain atom from which they branch, that is, atom 10 is the first

branch atom attached to chain atom 1, and atom 30 is the first branch atom attached to chain atom 3.

The Cartesian coordinates of the atoms are computed by the program from the equations in table I(a). Atom 1 is considered to be at the origin of a set of orthogonal axes and bond $r_{2,3}$ to be parallel to the x-axis. The general equations given in table I(a) will permit the extension of the program to accommodate more chain and branch atoms. Presently, the program can treat a nine-atom chain with two-atom branches from any of the first eight chain atoms.

The necessary input data are the appropriate bond lengths and angles according to the following assignment:

(1) The adjacent bond lengths are specified in order along the chain starting from atom 1 with bond $r_{1,2}$.

(2) The chain angles are specified in order along the chain beginning with angle ϕ_2 between the first and second bonds. The assigned angles are always on the same side, that is, the "inside," of the chain.

(3) For any particular branch, the bond lengths and branch angles are specified in order from the chain; all angles are taken on the same side of the branch, as indicated in figure 2(a-2).

Although $(\text{HBO}_2)_3$ is a closed cyclic structure, it is illustrated in figure 2(a-2) as an open chain with the bond length $r_{6,1}$ and angle ϕ_6 omitted. These parameters are unnecessary because the preceding bond lengths and angles have specified all the atom positions.

Other chain assignments are also possible for $(\text{HBO}_2)_3$, for example, an eight-atom chain with origin at the illustrated atom 11; the two branches would then be attached to the fifth and seventh chain atoms.

The equations for the atom coordinates do not permit branching from the last chain atom. If this error is made in the branch assignment, the computation stops and a suggested assignment is printed.

The input for the simplest chain, a diatomic molecule, consists only of the bond length.

Sample computations for planar molecules without and with branching are given in table II(a).

Type 2 - Pyramidal XY_3

Several molecules such as NH_3 , PH_3 , PCl_3 , and PF_3 are of the pyramidal XY_3 geometric type. The input for this type can be simplified to the X-Y bond length and the Y-X-Y angle.

A three-dimensional pyramidal molecule, oriented relative to a convenient set of orthogonal axes, is illustrated in figure 2(b). The figure also includes diagrams of the atom projections on the xy-, xz-, and yz-planes. The X atom is at the origin of the orthogonal axes and the Y_1 and the Y_2 atoms lie in the xz-plane. The z-axis bisects the Y_1 -X- Y_2 angle.

The equations for the Cartesian coordinates of the atoms illustrated in figure 2(b) are given in table I(b). Because of the selection of the coordinate system, these equations are simple functions of the X-Y bond length r and the Y-X-Y angle α .

A sample computation for a pyramidal XY_3 molecule is given in table II(b).

Type 3 - Tetrahedral XY_4

Molecules such as CCl_4 , $SiCl_4$, SiF_4 , and SiH_4 are of the tetrahedral XY_4 geometric type. The only input necessary for this type is the X-Y bond length.

A three-dimensional tetrahedral molecule, relative to a convenient set of orthogonal axes, is illustrated in figure 2(c). The figure also includes diagrams of the atom projections on the xy-, xz-, and yz-planes. The Y_1 , Y_2 , and Y_3 atoms lie in the xz-plane with Y_2 on the z-axis, and the X and Y_4 atoms lie on the y-axis.

The equations for the Cartesian coordinates of the atoms illustrated in figure 2(c) are given in table I(c). Because of the selection of the coordinate system, these equations are simple functions of the X-Y bond length r .

Sample computations for the tetrahedral XY_4 molecules are given in table II(c). The input for the P_4 computation involves a fictitious X atom, which is discussed in the section, SPECIAL TREATMENTS OF MOLECULES.

Type 4 - Other Nonplanar

Certain molecules have one or two atoms that are not in the same plane as the other atoms. Examples of these molecules are $(NaOH)_2$, S_2Cl_2 , and SF_4 . Their out-of-plane atoms are the two H atoms, one Cl atom, and two F atoms, respectively. Three-dimensional diagrams of these molecules, oriented relative to a convenient set of orthogonal axes, are sketched in figure 3. The input is specified by considering the out-of-plane atoms to be special branches from a planar chain. In addition to the bond lengths and angles required for the planar chain input, the out-of-plane-atom (hereinafter called O-P-A) bond length and angle are necessary.

In figure 2(d) a three-dimensional nonplanar molecule is illustrated with the atoms oriented relative to a convenient set of orthogonal axes. The figure also includes diagrams of the atom projections on the xy- and xz-planes. The

out-of-plane atom is specified according to the chain atom from which it branches; that is, out-of-plane atom 200 is attached to chain atom 2. A single-atom O-P-A branch is permitted only from chain atom 2 or 3. This restriction, however, is not severe, and many nonplanar molecules, including all those of interest in reference 2, can be described in this manner.

The Cartesian coordinates of the out-of-plane atoms illustrated in figure 2(d) are computed from the equations in table I(d). The O-P-A bond length $r_{i,k}$ is the distance between the out-of-plane atom and its adjacent chain atom. The O-P-A angle σ_i is formed by the out-of-plane atom, its adjacent chain atom, and one other chain atom, provided that these three atoms form a plane perpendicular to the plane of the chain. The intersection of these two planes is parallel to the x-axis and contains the line segment corresponding to bond length $r_{2,3}$. A plus or a minus sign indicates that the out-of-plane atom is above or below the plane of the chain.

Sample computations for other nonplanar molecules are given in table II(d). Special discussions of the input for the $(\text{NaOH})_2$ and the SF_4 computations are reserved for a later section, SPECIAL TREATMENTS OF MOLECULES.

Types 5 and 6 - General: Coordinates Read In

In these two types the atom coordinates may be read in either in the form $U + V \cos \alpha$ (Type 5) or directly as x, y, z (Type 6). When the orthogonal axes are judiciously chosen, the Cartesian coordinates of the atoms are usually simple functions of the bond lengths and angles. Types 5 and 6 are useful in the computation of principal moments of inertia for molecules not included in the other types and as checks on the other computations.

Examples of Type 5 and Type 6 computations are given in tables II(e) and (f), respectively. The molecules, oriented relative to the assigned coordinate system, are illustrated in figures 2(e) and (f).

SPECIAL TREATMENTS OF MOLECULES

A facility in visualization of structure is helpful in choosing the simplest way of treating a molecule. This is particularly the case with Type 4 - other nonplanar molecules. Occasionally the intuitive order of atoms along a chain is not the most convenient. For instance, the order of the chain atoms for $(\text{NaOH})_2$, illustrated in figure 3(a), was taken to be Na O O in order to accommodate the H atoms as O-P-A branches.

Certain molecules can be made to fit into a desired geometric-type classification by the invention of fictitious atoms of zero mass. The molecule SF_4 , shown in figure 3(c), was assigned a fictitious third chain atom in order to permit a planar branch and two O-P-A branches from the second chain atom. Thus the O-P-A plane is perpendicular to the plane of the chain.

Another example of a molecule for which a fictitious atom was invented is P₄. Ordinarily a Type 2 - pyramidal XY₃, this molecule can be treated as a Type 3 - tetrahedral XY₄. In this way, the available experimental data are more easily accommodated. The X atom is fictitious and the four Y atoms correspond to the P atoms. The required bond length input is the fictitious XY bond length, which is readily calculated from the experimental data. Since a fictitious atom has zero mass, it does not affect the principal moments of inertia.

Lewis Research Center

National Aeronautics and Space Administration
Cleveland, Ohio, November 13, 1963

REFERENCES

1. Hirschfelder, Joseph O.: Simple Method for Calculating Moments of Inertia. Jour. Chem. Phys., vol. 8, no. 5, May 1940, p. 431.
2. McBride, Bonnie J., Heimel, Sheldon, Ehlers, Janet G., and Gordon, Sanford: Thermodynamic Properties to 6000° K for 210 Substances Involving the First 18 Elements. NASA SP-3001, 1963.
3. Cameron, A. E., and Wichers, Edward: Report on the International Commission on Atomic Weights. Jour. Am. Chem. Soc., vol. 84, no. 22, Nov. 20, 1962, pp. 4175-4197.
4. Goldstein, Herbert.: Classical Mechanics. Addison-Wesley Pub. Co., Inc., 1959, pp. 143-184.
5. Greenstadt, John: The Determination of the Characteristic Roots of a Matrix by the Jacobi Method. Ch. 7 of Mathematical Methods for Digital Computers, Anthony Ralston and Herbert S. Wilf, eds., John Wiley & Sons, Inc., 1960, pp. 84-91.

TABLE I. - EQUATIONS FOR THE CARTESIAN COORDINATES OF THE ATOMS

(a) Type 1 - planar

Atom	x-coordinate, angstroms	y-coordinate, angstroms
1	0	0
2	$-r_{1,2} \cos \varphi_2$	$r_{1,2} \sin \varphi_2$
3	$x_2 + r_{2,3}$	y_2
4	$x_3 - r_{3,4} \cos \varphi_3$	$y_3 - r_{3,4} \sin \varphi_3$
5	$x_4 + r_{4,5} \cos(\varphi_3 + \varphi_4)$	$y_4 + r_{4,5} \sin(\varphi_3 + \varphi_4)$
^a 1	$x_{1-1} + (-1)^{1-1} r_{1-1,1} \cos\left(\sum_{k=3}^{1-1} \varphi_k\right)$	$y_{1-1} + (-1)^{1-1} r_{1-1,1} \sin\left(\sum_{k=3}^{1-1} \varphi_k\right)$
10	$-r_{1,10} \cos(\theta_1 - \varphi_2)$	$-r_{1,10} \sin(\theta_1 - \varphi_2)$
11	$x_{10} + r_{10,11} \cos(\theta_1 + \theta_{10} - \varphi_2)$	$y_{10} + r_{10,11} \sin(\theta_1 + \theta_{10} - \varphi_2)$
20	$x_2 + r_{2,20} \cos \theta_2$	$y_2 + r_{2,20} \sin \theta_2$
21	$x_{20} - r_{20,21} \cos(\theta_2 + \theta_{20})$	$y_{20} - r_{20,21} \sin(\theta_2 + \theta_{20})$
30	$x_3 - r_{3,30} \cos(\theta_3 + \varphi_3)$	$y_3 - r_{3,30} \sin(\theta_3 + \varphi_3)$
31	$x_{30} + r_{30,31} \cos(\theta_3 + \theta_{30} + \varphi_3)$	$y_{30} + r_{30,31} \sin(\theta_3 + \theta_{30} + \varphi_3)$
40	$x_4 + r_{4,40} \cos(\theta_4 + \varphi_3 + \varphi_4)$	$y_4 + r_{4,40} \sin(\theta_4 + \varphi_3 + \varphi_4)$
41	$x_{40} - r_{40,41} \cos(\theta_4 + \theta_{40} + \varphi_3 + \varphi_4)$	$y_{40} - r_{40,41} \sin(\theta_4 + \theta_{40} + \varphi_3 + \varphi_4)$
50	$x_5 - r_{5,50} \cos(\theta_5 + \varphi_3 + \varphi_4 + \varphi_5)$	$y_5 - r_{5,50} \sin(\theta_5 + \varphi_3 + \varphi_4 + \varphi_5)$
51	$x_{50} + r_{50,51} \cos(\theta_5 + \theta_{50} + \varphi_3 + \varphi_4 + \varphi_5)$	$y_{50} + r_{50,51} \sin(\theta_5 + \theta_{50} + \varphi_3 + \varphi_4 + \varphi_5)$
^b 101	$x_1 + (-1)^1 r_{1,101} \cos\left(\theta_1 + \sum_{k=3}^1 \varphi_k\right)$	$y_1 + (-1)^1 r_{1,101} \sin\left(\theta_1 + \sum_{k=3}^1 \varphi_k\right)$
^b 101+1	$x_{101} + (-1)^{1-1} r_{101,101+1} \cos\left(\theta_1 + \theta_{101} + \sum_{k=3}^1 \varphi_k\right)$	$y_{101} + (-1)^{1-1} r_{101,101+1} \sin\left(\theta_1 + \theta_{101} + \sum_{k=3}^1 \varphi_k\right)$

^aGeneral equations for extension of chain.^bGeneral equations for additional branches.

TABLE I. - Concluded. EQUATIONS FOR THE CARTESIAN
COORDINATES OF THE ATOMS

(b) Type 2 - pyramidal XY_3

Atom	x-coordinate, angstroms	y-coordinate, angstroms	z-coordinate, angstroms
X	0	0	0
Y_1	$r \sin \frac{\alpha}{2}$	0	$r \cos \frac{\alpha}{2}$
Y_2	$-r \sin \frac{\alpha}{2}$	0	$r \cos \frac{\alpha}{2}$
Y_3	0	$a \left[r^2 \left(1 - \frac{k^2}{4 \cos^2 \frac{\alpha}{2}} \right) \right]^{1/2}$	$\frac{rk}{2 \cos \frac{\alpha}{2}}$

(c) Type 3 - tetrahedral XY_4

X	0	$r/3$	0
Y_1	$r \frac{\sqrt{6}}{3}$	0	$r \frac{\sqrt{2}}{3}$
Y_2	0	0	$-2r \frac{\sqrt{2}}{3}$
Y_3	$-r \frac{\sqrt{6}}{3}$	0	$r \frac{\sqrt{2}}{3}$
Y_4	0	$r \frac{4}{3}$	0

(d) Type 4 - other nonplanar

200	$x_2 + r_{2,200} \cos \sigma_2$	y_2	$(\pm)r_{2,200} \sin \sigma_2$
300	$x_3 - r_{3,300} \cos \sigma_3$	y_3	$(\pm)r_{3,300} \sin \sigma_3$

$$a_k = 1 - 3 \sin^2 \frac{\alpha}{2} + \cos^2 \frac{\alpha}{2}.$$

TABLE II. - SAMPLE COMPUTATION OUTPUTS FOR PRINCIPAL MOMENTS OF INERTIA

(a) Type 1

(a-1) HBO_2

PLANAR WITH 4 CHAIN ATOMS AND 0 BRANCHES. WHITE JCP V32 P488 FEB 1960.

NO.	CHAIN ATOM	BOND LENGTH	ANGLE	MOLECULAR WEIGHT	X	Y	Z
1	H	1.0000		1.00799999	0	0	0
2	O	1.3400	120.00	16.0000000	0.49999996	0.86602541	0
3	B	1.2000	180.00	10.8199999	1.83999993	0.86602541	0
4	O			16.0000000	3.03999990	0.86602537	0

THE GENERATED SYMMETRIC MATRIX IS

I				I
I	0.1226276	-0.2531333	0	I
I				I
I	-0.2531333	9.0980780	0	I
I				I
I	0	0	9.2207053	I
I				I

THE EIGENVALUES, LAMBDA, OF THE MATRIX ARE THE PRINCIPAL MOMENTS OF INERTIA (G-SQ CM)

LAMBDA(1)*1.0E+39 = 0.11549414
 LAMBDA(2)*1.0E+39 = 9.10521126
 LAMBDA(3)*1.0E+39 = 9.22070527

PRODUCT LAMBDA*1.0E+117 = 9.69648027

THE GENERATED MATRICES FROM THE JACOBI METHOD ARE AS FOLLOWS,

THE DIAGONAL MATRIX (D)

I				I
I	0.11549414	0	0	I
I				I
I	0	9.10521126	0	I
I				I
I	0	0	9.22070527	I
I				I

THE EIGENVECTOR MATRIX (S)

I				I
I	0.99960316	-0.28169271E-01	0	I
I				I
I	0.28169271E-01	0.99960316	0	I
I				I
I	0	-0	1.00000000	I
I				I

THE MATRIX (S)*(D)*(S TRANSPOSE), WHICH SHOULD EQUAL THE ORIGINAL MATRIX IS,

I				I
I	0.12262755	-0.25313329	0	I
I				I
I	-0.25313329	9.09807754	0	I
I				I
I	0	0	9.22070527	I
I				I

TABLE II. - Continued. SAMPLE COMPUTATION OUTPUTS FOR PRINCIPAL MOMENTS OF INERTIA

(a) Concluded. Type 1

(a-2) $(\text{HBO}_2)_3$

PLANAR WITH 6 CHAIN ATOMS AND 3 BRANCHES. WHITE JCP V32 P488 FEB. 1960 B-D 1.36A, O-H 1.A, CYCLIC

NO.	CHAIN ATOM	BOND LENGTH	ANGLE	MOLECULAR WEIGHT	X	Y	Z
1	B	1.3600		10.8199999	0	0	0
2	O	1.3600	120.00	16.0000000	0.67999993	1.17779455	0
3	B	1.3600	120.00	10.8199999	2.03999990	1.17779455	0
4	O	1.3600	120.00	16.0000000	2.71999982	-0	0
5	B	1.3600	120.00	10.8199999	2.03999972	-1.17779449	0
6	O	1.3600		16.0000000	0.67999974	-1.17779440	0

BRANCH NO. 1, ATTACHED TO CHAIN ATOM B, NUMBER 1

NO.	ATOM	BOND LENGTH	ANGLE				
7	O	1.3600	120.00	16.0000000	-1.35999998	-0	0
8	H	1.0000	240.00	1.00799999	-1.86000004	-0.86602537	0

BRANCH NO. 2, ATTACHED TO CHAIN ATOM B, NUMBER 3

NO.	ATOM	BOND LENGTH	ANGLE				
9	O	1.3600	120.00	16.0000000	2.71999997	2.35558903	0
10	H	1.0000	240.00	1.00799999	2.22000015	3.22161448	0

BRANCH NO. 3, ATTACHED TO CHAIN ATOM B, NUMBER 5

NO.	ATOM	BOND LENGTH	ANGLE				
11	O	1.3600	120.00	16.0000000	2.71999946	-2.35558909	0
12	H	1.0000	240.00	1.00799999	3.71999943	-2.35558921	0

THE GENERATED SYMMETRIC MATRIX IS

I				I
I	44.624305	-0.3702343E-05	0	I
I				I
I	-0.3702343E-05	44.624300	0	I
I				I
I	0	0	89.248606	I
I				I

THE EIGENVALUES, LAMBDA, OF THE MATRIX ARE THE PRINCIPAL MOMENTS OF INERTIA (G-SQ CM)

LAMBDA(1)*1.0E+39 = 44.6243057
 LAMBDA(2)*1.0E+39 = 44.6242971
 LAMBDA(3)*1.0E+39 = 89.2486057

PRODUCT LAMBDA*1.0E+117 = 177723.270

THE GENERATED MATRICES FROM THE JACOBI METHOD ARE AS FOLLOWS,

THE DIAGONAL MATRIX (D)

I				I
I	44.6243057	0	0	I
I				I
I	0	44.6242971	0	I
I				I
I	0	0	89.2486057	I
I				I

THE EIGENVECTOR MATRIX (S)

I				I
I	0.86644093	0.49927959	0	I
I				I
I	-0.49927959	0.86644093	0	I
I				I
I	0	0	1.00000000	I
I				I

THE MATRIX (S)*(D)*(S TRANSPOSE), WHICH SHOULD EQUAL THE ORIGINAL MATRIX IS,

I				I
I	44.6243029	-0.35762787E-05	0	I
I				I
I	-0.40531158E-05	44.6242981	0	I
I				I
I	0	0	89.2486057	I
I				I

TABLE II. - Continued. SAMPLE COMPUTATION OUTPUTS FOR PRINCIPAL MOMENTS OF INERTIA

(b) Type 2 - PCl_3

PYRAMIDAL WITH BOND LENGTH= 2.04300 , AND ANGLE= 100.10
KISLIUK JCP V18 P1109 AUG 1950.

NO.	ATOM	MOLECULAR WEIGHT	X	Y	Z
1	P	30.9749999	0	0	0
2	CL	35.4569998	1.56617413	0	1.31184886
3	CL	35.4569998	-1.56617413	0	1.31184886
4	CL	35.4569998	0	1.96533303	-0.55795603

THE GENERATED SYMMETRIC MATRIX IS

1	32.476808	0	0
1	0	44.488149	12.625000
1	0	12.625000	45.746817

THE EIGENVALUES, LAMBDA, OF THE MATRIX ARE THE
PRINCIPAL MOMENTS OF INERTIA (G-SQ CM)

LAMBDA(1)=1.0E+39 = 32.4768076
LAMBDA(2)=1.0E+39 = 32.4768052
LAMBDA(3)=1.0E+39 = 57.7581563

PRODUCT LAMBDA=1.0E+117 = 60920.0078

THE GENERATED MATRICES FROM THE JACOBI METHOD ARE AS FOLLOWS.

THE DIAGONAL MATRIX (D)

1	32.4768076	0	0
1	0	32.4768052	0
1	0	0	57.7581563

THE EIGENVECTOR MATRIX (S)

1	1.00000000	0	0
1	0	0.72449513	0.68927990
1	0	-0.68927990	0.72449513

THE MATRIX (S)*(D)*(S TRANSPOSE), WHICH
SHOULD EQUAL THE ORIGINAL MATRIX IS.

1	32.4768076	0	0
1	0	44.4881449	12.6249990
1	0	12.6249990	45.7468128

TABLE II. - Continued. SAMPLE COMPUTATION OUTPUTS FOR PRINCIPAL MOMENTS OF INERTIA

(c) Type 3

(c-1) CCl_4

TETRAHEDRAL-REGULAR WITH BOND LENGTH= 1.7600 BARTELL, JCP V23 P1854 OCT 1955.

NO.	ATOM	MOLECULAR WEIGHT	X	Y	Z
1	C	12.0109999	0	0.58666666	0
2	CL	35.4569998	1.43703395	0	0.82967194
3	CL	35.4569998	0	0	-1.65934388
4	CL	35.4569998	-1.43703395	0	0.82967194
5	CL	35.4569998	0	2.34666663	0

THE GENERATED SYMMETRIC MATRIX IS

I			
I	48.625861	0	0
I			
I	0	48.625861	0
I			
I	0	0	48.625861
I			

THE EIGENVALUES, LAMBDAS, OF THE MATRIX ARE THE PRINCIPAL MOMENTS OF INERTIA (G-SQ CM)

LAMBDA(1)=1.0E+39 = 48.6258612
LAMBDA(2)=1.0E+39 = 48.6258607
LAMBDA(3)=1.0E+39 = 48.6258612

PRODUCT LAMBDA=1.0E+117 = 114974.601

THE GENERATED MATRICES FROM THE JACOBI METHOD ARE AS FOLLOWS,

THE DIAGONAL MATRIX (D)

I			
I	48.6258612	0	0
I			
I	0	48.6258607	0
I			
I	0	0	48.6258612
I			

THE EIGENVECTOR MATRIX (S)

I			
I	1.00000000	0	0
I			
I	0	1.00000000	0
I			
I	0	0	1.00000000
I			

THE MATRIX (S)*(D)*(S TRANSPOSE), WHICH SHOULD EQUAL THE ORIGINAL MATRIX IS,

I			
I	48.6258612	0	0
I			
I	0	48.6258607	0
I			
I	0	0	48.6258612
I			

TABLE II. - Continued. SAMPLE COMPUTATION OUTPUTS FOR PRINCIPAL MOMENTS OF INERTIA

(c) Concluded. Type 3

(c-2) P_4

TETRAHEDRAL-REGULAR WITH BOND LENGTH= 1.3533 MAXWELL JCP V3 P699 NOV 1935.

NO.	ATOM	MOLECULAR WEIGHT	X	Y	Z
1		0	0	0.45111433	0
2	P	30.9749999	1.10499991	0	0.63797200
3	P	30.9749999	0	0	-1.27594399
4	P	30.9749999	-1.10499991	0	0.63797200
5	P	30.9749999	0	1.80445731	0

THE GENERATED SYMMETRIC MATRIX IS

I				I
I	25.116959	0	0	I
I				I
I	0	25.116959	0	I
I				I
I	0	0	25.116960	I
I				I

THE EIGENVALUES, LAMBDA, OF THE MATRIX ARE THE PRINCIPAL MOMENTS OF INERTIA (G-SQ CM)

LAMBDA(1)*1.0E+39 = 25.1169593
 LAMBDA(2)*1.0E+39 = 25.1169593
 LAMBDA(3)*1.0E+39 = 25.1169596

PRODUCT LAMBDA*1.0E+117 = 15845.3263

THE GENERATED MATRICES FROM THE JACOBI METHOD ARE AS FOLLOWS,

THE DIAGONAL MATRIX (D)

I				I
I	25.1169593	0	0	I
I				I
I	0	25.1169593	0	I
I				I
I	0	0	25.1169596	I
I				I

THE EIGENVECTOR MATRIX (S)

I				I				I
I	1.00000000	0	0	I				I
I				I				I
I	0	1.00000000	0	I				I
I				I				I
I	0	0	1.00000000	I				I
I				I				I

THE MATRIX (S)*(D)*(S TRANSPOSE), WHICH SHOULD EQUAL THE ORIGINAL MATRIX IS,

I				I
I	25.1169593	0	0	I
I				I
I	0	25.1169593	0	I
I				I
I	0	0	25.1169596	I
I				I

TABLE II. - Continued. SAMPLE COMPUTATION OUTPUTS FOR PRINCIPAL MOMENTS OF INERTIA

(d) Type 4

(d-1) (NaOH)₂

NON-PLANAR WITH 3 CHAIN ATOMS, 1 PLANAR BRANCHES, AND 2 OUT-OF-PLANE ATOMS.

ESTIMATES FROM NAOH AND LIOH, (LIOH)₂.

NO.	CHAIN ATOM	BOND LENGTH	ANGLE	MOLECULAR WEIGHT	X	Y	Z
1	NA			22.9909999	0	0	0
2	O	2.2500	40.00	16.0000000	-1.72360000	1.44627209	0
3	O	3.4472		16.0000000	1.72359994	1.44627209	0
BRANCH NO. 1, ATTACHED TO CHAIN ATOM O , NUMBER 2							
NO. 4	ATOM NA	BOND LENGTH 2.2500	ANGLE 40.00	22.9909999	0	2.89254418	0
O-P-A BRANCH NO. 1, ATTACHED TO CHAIN ATOM O , NUMBER 2							
NO. 5	ATOM H	BOND LENGTH 0.9600	O-P-A ANGLE 120.00	1.00799999	-2.20359993	1.44627209	0.83138438
O-P-A BRANCH NO. 2, ATTACHED TO CHAIN ATOM O , NUMBER 3							
NO. 6	ATOM H	BOND LENGTH 0.9600	O-P-A ANGLE 120.00	1.00799999	2.20359987	1.44627209	-0.83138438

THE GENERATED SYMMETRIC MATRIX IS

I	I	I	I	I
I	16.199666	-0.4701138E-07	0.6131919	I
I	I	I	I	I
I	I	-0.4701138E-07	17.639793	-0
I	I	I	I	I
I	I	I	I	I
I	I	I	I	I
I	I	0.6131919	-0	33.376763
I	I	I	I	I

THE EIGENVALUES, LAMBDA, OF THE MATRIX ARE THE PRINCIPAL MOMENTS OF INERTIA (G-SQ CM)

LAMBDA(1)=1.0E+39 = 16.1778021
 LAMBDA(2)=1.0E+39 = 17.6397927
 LAMBDA(3)=1.0E+39 = 33.3986244

PRODUCT LAMBDA*1.0E+117 = 9531.06799

THE GENERATED MATRICES FROM THE JACOBI METHOD ARE AS FOLLOWS,

THE DIAGONAL MATRIX (D)

I	I	I	I	I
I	16.1778021	0.72469895E-18	0.55511151E-16	I
I	I	I	I	I
I	I	0.72469895E-18	-0.27755576E-16	I
I	I	I	I	I
I	I	I	I	I
I	I	I	I	I
I	I	0.55511151E-16	-0.27755576E-16	33.3986244
I	I	I	I	I

THE EIGENVECTOR MATRIX (S)

I	I	I	I	I
I	0.99936502	-0.32111128E-07	0.35630208E-01	I
I	I	I	I	I
I	I	0.32135320E-07	0.99999999	-0.10629120E-09
I	I	I	I	I
I	I	I	I	I
I	I	I	I	I
I	I	-0.35630208E-01	0.12512118E-08	0.99936503
I	I	I	I	I

THE MATRIX (S)*(D)*(S TRANSPOSE), WHICH SHOULD EQUAL THE ORIGINAL MATRIX IS,

I	I	I	I	I
I	16.1996632	-0.47011378E-07	0.61319189	I
I	I	I	I	I
I	I	-0.47011378E-07	17.6397922	0.47184478E-15
I	I	I	I	I
I	I	I	I	I
I	I	I	I	I
I	I	0.61319188	-0.16653345E-15	33.3767610
I	I	I	I	I

TABLE II. - Continued. SAMPLE COMPUTATION OUTPUTS FOR PRINCIPAL MOMENTS OF INERTIA

(d) Continued. Type 4

(d-2) S_2Cl_2

NON-PLANAR WITH 3 CHAIN ATOMS, 0 PLANAR BRANCHES, AND 1 OUT-OF-PLANE ATOMS.

BOWEN, CHEM. SOC., SPECIAL PUBL. NO. 11, 1958.

NO.	CHAIN ATOM	BOND LENGTH	ANGLE	MOLECULAR WEIGHT	X	Y	Z
1	CL	1.9900		35.4569998	0	0	0
2	S	2.0500	104.00	32.0660000	0.48142453	1.93088849	0
3	S			32.0660000	2.53142446	1.93088849	0

O-P-A BRANCH NO. 1, ATTACHED TO CHAIN ATOM S, NUMBER 3

NO.	ATOM	BOND LENGTH	O-P-A ANGLE	MOLECULAR WEIGHT	X	Y	Z
4	CL	1.9900	104.00	35.4569998	3.01284897	1.93088849	-1.93088849

THE GENERATED SYMMETRIC MATRIX IS

I				I
I	32.370342	-17.122919	17.122920	I
I				I
I	-17.122919	54.089292	5.7624598	I
I				I
I	17.122920	5.7624598	54.089291	I
I				I

THE EIGENVALUES, LAMBDA, OF THE MATRIX ARE THE PRINCIPAL MOMENTS OF INERTIA (G-SQ CM)

LAMBDA(1)*1.0E+39 = 14.8526815
 LAMBDA(2)*1.0E+39 = 65.8444834
 LAMBDA(3)*1.0E+39 = 59.8517456

PRODUCT LAMBDA*1.0E+117 = 58533.0405

THE GENERATED MATRICES FROM THE JACOBI METHOD ARE AS FOLLOWS,

THE DIAGONAL MATRIX (D)

I				I
I	14.8526815	0	0.82881942E-12	I
I				I
I	0	65.8444834	0	I
I				I
I	0.82881942E-12	0	59.8517456	I
I				I

THE EIGENVECTOR MATRIX (S)

I				I
I	0.81022295	-0.58612171	-0.24990641E-08	I
I				I
I	0.41445062	0.57291412	0.70710676	I
I				I
I	-0.41445065	-0.57291416	0.70710672	I
I				I

THE MATRIX (S)*(D)*(S TRANSPOSE), WHICH SHOULD EQUAL THE ORIGINAL MATRIX IS,

I				I
I	32.3703346	-17.1229138	17.1229153	I
I				I
I	-17.1229141	54.0892787	5.76245952	I
I				I
I	17.1229153	5.76245928	54.0892787	I
I				I

TABLE II. - Continued. SAMPLE COMPUTATION OUTPUTS FOR PRINCIPAL MOMENTS OF INERTIA

(d) Concluded. Type 4

(d-3) SF₄

NON-PLANAR WITH 3 CHAIN ATOMS, 1 PLANAR BRANCHES, AND 2 OUT-OF-PLANE ATOMS.

DODD TRANS FAR SOC V52 P1052 1956. FICTITIOUS THIRD ATOM.

NO.	CHAIN ATOM	BOND LENGTH	ANGLE	MOLECULAR WEIGHT	X	Y	Z
1	F	1.5800		19.0000000	0	0	0
2	S	0.7900	60.00	32.0660000	-0.79000001	1.36832011	0
3				0	-0.14901161E-07	1.36832011	0
BRANCH NO. 1, ATTACHED TO CHAIN ATOM S, NUMBER 2							
NO. 4	ATOM F	BOND LENGTH 1.5800	ANGLE 60.00	19.0000000	0	2.73664021	0
O-P-A BRANCH NO. 1, ATTACHED TO CHAIN ATOM S, NUMBER 2							
NO. 5	ATOM F	BOND LENGTH 1.5800	O-P-A ANGLE 110.00	19.0000000	-1.33039176	1.36832011	1.48471433
O-P-A BRANCH NO. 2, ATTACHED TO CHAIN ATOM S, NUMBER 2							
NO. 6	ATOM F	BOND LENGTH 1.5800	O-P-A ANGLE 110.00	19.0000000	-1.33039176	1.36832011	-1.48471433

THE GENERATED SYMMETRIC MATRIX IS

I				I
I	25.719417	-0	0	I
I				I
I				I
I	-0	19.548759	-0	I
I				I
I				I
I	0	-0	17.453714	I
I				I

THE EIGENVALUES, LAMBDA, OF THE MATRIX ARE THE PRINCIPAL MOMENTS OF INERTIA (G-SQ CM)

LAMBDA(1)*1.0E+39 = 25.7194171
 LAMBDA(2)*1.0E+39 = 19.5487585
 LAMBDA(3)*1.0E+39 = 17.4537139

PRODUCT LAMBDA*1.0E+117 = 8775.42480

THE GENERATED MATRICES FROM THE JACOBI METHOD ARE AS FOLLOWS,

THE DIAGONAL MATRIX (D)

I				I
I	25.7194171	-0	0	I
I				I
I	-0	19.5487585	-0	I
I				I
I	0	-0	17.4537139	I
I				I

THE EIGENVECTOR MATRIX (S)

I				I
I	1.00000000	0	0	I
I				I
I	0	1.00000000	0	I
I				I
I	0	0	1.00000000	I
I				I

THE MATRIX (S)*(D)*(S TRANSPOSE), WHICH SHOULD EQUAL THE ORIGINAL MATRIX IS,

I				I
I	25.7194171	0	0	I
I				I
I	0	19.5487585	0	I
I				I
I	0	0	17.4537139	I
I				I

TABLE II. - Continued. SAMPLE COMPUTATION OUTPUTS FOR PRINCIPAL MOMENTS OF INERTIA

(e) Type 5 - SOCl_2

COORDINATES READ IN-- U+VCOS(ALPHA) PALMER JACS V60 P2360 1938. S=0 1.45A, S-CL 2.07A, O-S-CL 106, CL-S-CL 114.

U,V,ALF INPUT			X			Y			Z		
ATOM	U	V	ALPHA	U	V	ALPHA	U	V	ALPHA		
S	-0.	0.	-0.	-0.	-0.	-0.	-0.	-0.	-0.		
O	-0.	-0.7338	-0.	-0.	1.2506	-0.	-0.	-0.	-0.		
CL	-0.	2.0700	57.0000	-0.	-0.	-0.	-0.	2.0700	33.0000		
CL	-0.	2.0700	57.0000	-0.	-0.	-0.	-0.	-2.0700	33.0000		

NO.	ATOM	MOLECULAR WEIGHT	X	Y	Z
1	S	32.0660000	0	-0	-0
2	O	16.0000000	-0.73383599	1.25059398	-0
3	CL	35.4569998	1.12740281	-0	1.73604809
4	CL	35.4569998	1.12740281	-0	-1.73604809

THE GENERATED SYMMETRIC MATRIX IS

I			
I	39.079384	4.3422745	-0
I			
I	4.3422745	45.386801	-0
I			
I	-0	-0	13.499138
I			

THE EIGENVALUES, LAMBDA, OF THE MATRIX ARE THE PRINCIPAL MOMENTS OF INERTIA (G-SQ CM)

LAMBDA(1)*1.0E+39 = 36.8664145
 LAMBDA(2)*1.0E+39 = 47.5997691
 LAMBDA(3)*1.0E+39 = 13.4991376

PRODUCT LAMBDA*1.0E+117 = 23688.7295

THE GENERATED MATRICES FROM THE JACOBI METHOD ARE AS FOLLOWS,

THE DIAGONAL MATRIX (D)

I			
I	36.8664145	0	0
I			
I	0	47.5997691	0
I			
I	0	0	13.4991376
I			

THE EIGENVECTOR MATRIX (S)

I			
I	0.89096754	0.45406698	0
I			
I	-0.45406698	0.89096754	0
I			
I	0	0	1.00000000
I			

THE MATRIX (S)*(D)*(S TRANSPOSE), WHICH SHOULD EQUAL THE ORIGINAL MATRIX IS.

I			
I	39.0793824	4.34227443	0
I			
I	4.34227455	45.3867989	0
I			
I	0	0	13.4991376
I			

TABLE II. - Concluded. SAMPLE COMPUTATION OUTPUTS FOR PRINCIPAL MOMENTS OF INERTIA
(r) Type 6 - S₂Cl₂

COORDINATES READ IN-- X, Y, Z BOWEN, CHEM. SOC., SPECIAL PUBL. NO. 11, 1958.

NO.	ATOM	MOLECULAR WEIGHT	X	Y	Z
1	S	32.0660000	0	0	0
2	S	32.0660000	2.04999998	0	0
3	CL	35.4569998	-0.48142456	1.93088858	0
4	CL	35.4569998	2.53142080	0	1.93088849

THE GENERATED SYMMETRIC MATRIX IS

I				I
I	32.370343	17.122910	-17.122889	I
I				I
I	17.122910	54.089228	5.7624599	I
I				I
I	-17.122889	5.7624599	54.089231	I
I				I

THE EIGENVALUES, LAMBDA, OF THE MATRIX ARE THE
PRINCIPAL MOMENTS OF INERTIA (G-SQ CM)

LAMBDA(1)*1.0E+39 = 14.8526877
LAMBDA(2)*1.0E+39 = 65.8444166
LAMBDA(3)*1.0E+39 = 59.8516831

PRODUCT LAMBDA*1.0E+117 = 58532.9443

THE GENERATED MATRICES FROM THE JACOBI METHOD ARE AS FOLLOWS.

THE DIAGONAL MATRIX (D)

I				I
I	14.8526877	-0.88817842E-15	-0.46649288E-09	I
I				I
I	-0.88817842E-15	65.8444166	0.45474811E-12	I
I				I
I	-0.46649288E-09	0.45474811E-12	59.8516831	I
I				I

THE EIGENVECTOR MATRIX (S)

I				I
I	0.81022272	0.58612205	-0.49451341E-06	I
I				I
I	-0.41445107	0.57291483	0.70710592	I
I				I
I	0.41445067	-0.57291310	0.70710757	I
I				I

THE MATRIX (S)*(D)*(S TRANSPOSE), WHICH
SHOULD EQUAL THE ORIGINAL MATRIX IS,

I				I
I	32.3703361	17.1229050	-17.1228845	I
I				I
I	17.1229053	54.0892148	5.76245952	I
I				I
I	-17.1228848	5.76245928	54.0892172	I
I				I

END OF FILE TAPE A 2

TABLE III. - CODE LISTING

```

C MOMENTS OF INERTIA                                MOM 001
C PROGRAM COMPUTES PRINCIPAL MOMENTS OF INERTIA FOR THE FOLLOWING TYPE MOM 002
C OF SUBSTANCES                                     MOM 003
C   NTYPE=1    PLANAR                                EFN 100    MOM 004
C   NTYPE=2    PYRAMIDAL                             EFN 200    MOM 005
C   NTYPE=3    TETRAHEDRAL                           EFN 300    MOM 006
C   NTYPE=4    NON-PLANAR WITH OUT-OF-PLANE ATOMS, (O-P-A) EFN 400    MOM 007
C   NTYPE=5    READ DATA (U, V, AND ALPHA), FOR EACH X,Y,Z EFN 500    MOM 008
C   NTYPE=6    READ COORDINATES OF EACH ATOM, X, Y, Z   EFN 600    MOM 009
C                                                     MOM 010
C                                                     MOM 011
C                                                     MOM 012
C PERMANENT INPUT                                    MOM 013
C                                                     MOM 014
C CARD 1  CC 1-3    NO. OF ENTRIES IN MOLECULAR WEIGHT TABLES.    INTEGER MOM 015
C          CC 4-11  EN, AVOGADRO NO. *1.0E-23                     FLOAT  MOM 016
C          CC 12-19, 20-27, ETC. MOLECULAR WEIGHTS OF ATOMS,      MOM 017
C                               IN THE ORDER TO BE LISTED ON      MOM 018
C                               NEXT CARD. OVERFLOW ONTO           MOM 019
C                               ADDITIONAL CARDS IF NECESSARY,     MOM 020
C                               STARTING IN CC1 WITH 8CC/ENTRY,    MOM 021
C                               AND MAXIMUM OF 10 WEIGHTS / CARD.  MOM 022
C CARD 2  NAMES OF ATOMS FOR WHICH MOLECULAR WEIGHTS HAVE BEEN    MOM 023
C        READ IN. 2 CC / NAME, LEFT ADJUSTED, MAXIMUM 40 ATOM NAMES / CARD. MOM 024
C        OVERFLOW ONTO ADDITIONAL CARDS IF NECESSARY.            MOM 025
C                                                     MOM 026
C                                                     MOM 027
C INPUT FORMAT FOR EACH SUBSTANCE.                      MOM 028
C CARD 3  CC 1-78 COMMENTS CARD.                          ALPH  MOM 029
C          CC 79-80  THE 2 CHARACTERS C1.                  MOM 030
C                                                     MOM 031
C                                                     MOM 032
C CARD 4  CC 1-3    NUMBER OF ATOMS.                        INTEGER MOM 033
C          CC 4-6    TYPE OF MOLECULE, NTYPE=1-6 AS LISTED ABOVE. INTEGR MOM 034
C          CC 7-30   NAME OF SUBSTANCE, EXAMPLE, BE(OH)2 .      ALPH  MOM 035
C          CC 31-78  NAME OF EACH ATOM, IN DESIRED ORDER,      MOM 036
C                    2CC/ATOM NAME, LEFT ADJUSTED,            MOM 037
C                    MAXIMUM 24 ATOMS. EXAMPLE, H O BEO H .    MOM 038
C                                                     MOM 039
C THE REMAINING INPUT DEPENDS ON THE TYPE.              MOM 040
C   BOND LENGTHS - ANGSTROMS.                             MOM 041
C   ANGLES        - DEGREES.                              MOM 042
C                                                     MOM 043
C                                                     MOM 044
C NTYPE=1 , PLANAR.                                       MOM 045
C                                                     MOM 046
C CARD 5  CC 1-3    NUMBER OF CHAIN ATOMS = N.              INTEGER MOM 047
C          CC 4-6    NUMBER OF BRANCHES =M.                 INTEGER MOM 048
C                                                     MOM 049
C CARD 6  CC 1-10, 11-20, ETC., (N-1) BOND LENGTHS,        FLOAT MOM 050
C          (N-2) ANGLES,                                     FLOAT MOM 051
C          OVERFLOW ONTO NEXT CARD (6A) IF NECESSARY (MAX 8 WDS./CD) MOM 052
C          ALL BOND LENGTHS ARE LISTED FIRST, THEN ALL ANGLES. MOM 053
C                                                     MOM 054
C IF THERE ARE ANY PLANAR BRANCHES,                      MOM 055
C CARD 7  CC 1-3    NO. OF CHAIN ATOM TO WHICH BRANCH IS ATTACHED. INT MOM 056

```


TABLE III. - Continued. CODE LISTING

C	CC 4-6 NO. OF ATOMS ON THIS BRANCH (N).	INTEGER	MOM	057
C	CARD 8 CC 1-10, 11-20, ETC., N BOND LENGTHS, N ANGLES.	FLOAT	MOM	058
C			MOM	059
C	REPEAT CARDS 7 AND 8 FOR EACH PLANAR BRANCH.		MOM	060
C			MOM	061
C			MOM	062
C	CNTYPE=2 , PYRAMIDAL.		MOM	063
C			MOM	064
C	CARD 5 CC 1-10 BOND LENGTH.	FLOAT	MOM	065
C	CC 11-20 ANGLE.	FLOAT	MOM	066
C			MOM	067
C			MOM	068
C	CNTYPE=3 , TETRAHEDRAL.		MOM	069
C			MOM	070
C	CARD 5 CC 1-10 BOND LENGTH.	FLOAT	MOM	071
C			MOM	072
C			MOM	073
C	CNTYPE=4 , NON-PLANAR WITH OUT-OF-PLANE ATOMS, (O-P-A).		MOM	074
C			MOM	075
C	CARD 5 CC 1-3 NO. OF CHAIN ATOMS.	INTEGER	MOM	076
C	CC 4-6 NO. OF PLANAR BRANCHES.	INTEGER	MOM	077
C	CC 7-9 NO. OF O-P-A BRANCHES.	INTEGER	MOM	078
C			MOM	079
C	INSTRUCTIONS FOR CARDS 6, 7, AND 8, SAME AS FOR PLANAR, NTYPE=1.		MOM	080
C			MOM	081
C	CARD 9 CC 1-3 NO. OF CHAIN ATOM TO WHICH O-P-A BRANCH	INTEGER	MOM	082
C	IS ATTACHED.		MOM	083
C	CC 11-20 BOND LENGTH.	FLOAT	MOM	084
C	CC 21-30 BOND ANGLE.	FLOAT	MOM	085
C	CC 31-40 SIGN (+1.0 , OR -1.0), TO INDICATE ABOVE	FLOAT	MOM	086
C	OR BELOW THE PLANE.		MOM	087
C	REPEAT CARD 9 FOR EACH O-P-A BRANCH.		MOM	088
C			MOM	089
C			MOM	090
C	CNTYPE=5 , READ DATA (U, V, ALPHA), IN THE FORM U+VCOS(ALPHA),		MOM	091
C	FOR X, Y, Z, OF EACH ATOM.		MOM	092
C			MOM	093
C	CARDS CC 1- 8 U , TO CALCULATE X OF GIVEN ATOM.	FLOAT	MOM	094
C	CC 9-16 V , TO CALCULATE X OF GIVEN ATOM.	FLOAT	MOM	095
C	CC 17-24 ALPHA, TO CALCULATE X OF GIVEN ATOM.	FLOAT	MOM	096
C	CC 25-32 U , TO CALCULATE Y OF GIVEN ATOM.	FLOAT	MOM	097
C	CC 33-40 V , TO CALCULATE Y OF GIVEN ATOM.	FLOAT	MOM	098
C	CC 41-48 ALPHA, TO CALCULATE Y OF GIVEN ATOM.	FLOAT	MOM	099
C	CC 49-56 U , TO CALCULATE Z OF GIVEN ATOM.	FLOAT	MOM	100
C	CC 57-64 V , TO CALCULATE Z OF GIVEN ATOM.	FLOAT	MOM	101
C	CC 65-72 ALPHA, TO CALCULATE Z OF GIVEN ATOM.	FLOAT	MOM	102
C	REPEAT CARD 5 FOR EACH ATOM OF SUBSTANCE, N ATOMS=N CARDS.		MOM	103
C			MOM	104
C			MOM	105
C	CNTYPE=6, COORDINATES READ IN, X, Y, Z.		MOM	106
C	CARD 5 CC 1-10,11-20,ETC., X OF EACH ATOM IN ORDER LISTED.	FLOAT	MOM	107
C	CARD 6 CC 1-10,11-20,ETC., Y OF EACH ATOM IN ORDER LISTED.	FLOAT	MOM	108
C	CARD 7 CC 1-10,11-20,ETC., Z OF EACH ATOM IN ORDER LISTED.	FLOAT	MOM	109
C	COMMON A,S,SD		MOM	110
C	EQUIVALENCE (CHECK,NCHECK),(NCOM,COMENT(14))		MOM	111
C	EQUIVALENCE (A,D)		MOM	112

TABLE III. - Continued. CODE LISTING

```

      DIMENSION A(3,3),D(3,3),S(3,3),SD(3,3)          MOM 113
      DIMENSION SUBS(4),COMENT(14),MOL(26),X(26),Y(26),Z(26),EMN(26),EM( MOM 114
126),WGHT(105),NAME(105),BOND(25),ANG(24),V(3),ANGBR(2) MOM 115
C                                     MOM 116
C SET IT1 = TO NUMBER OF INPUT TAPE                     MOM 117
C                                     MOM 118
C SET IT2 = TO NUMBER OF OUTPUT TAPE                   MOM 119
C                                     MOM 120
      IT1=7                                              MOM 121
      IT2=6                                              MOM 122
C                                     MOM 123
      RAD=0.0174532925                                  MOM 124
C                                     MOM 125
C READ PERMANENT DATA.                                MOM 126
C                                     MOM 127
      READ INPUT TAPE IT1,1002,NTABL,EN,(WGHT(J),J=1,NTABL) MOM 128
      NTABL=NTABL+1                                     MOM 129
      WGHT(NTABL)=0.                                    MOM 130
      READ INPUT TAPE IT1,1006,(NAME(J),J=1,NTABL)      MOM 131
C                                     MOM 132
C READ INPUT CARDS 3 AND 4, INPUT FOR EACH SUBSTANCE.  MOM 133
C                                     MOM 134
      900 READ INPUT TAPE IT1,1003,(COMENT(J),J=1,14)   MOM 135
      901 READ INPUT TAPE IT1,1000,NATMS,NTYPE,(SUBS(J),J=1,4), MOM 136
      1(MOL(J),J=1,NATMS)                               MOM 137
      WRITE OUTPUT TAPE IT2,2015,(SUBS(J),J=1,4)        MOM 138
      SWITCH=0.                                         MOM 139
C                                     MOM 140
C GET MOLECULAR WEIGHT OF EACH ATOM                    MOM 141
C                                     MOM 142
      DO 801 J=1,NATMS                                  MOM 143
      IF (MOL(J)-NAMOLD)802,803,802                     MOM 144
803 EMN(J)=WTOLD                                         MOM 145
      GO TO 40                                           MOM 146
802 DO 804 K=1,NTABL                                    MOM 147
      IF (MOL(J)-NAME(K))804,805,804                    MOM 148
805 KK=K                                                 MOM 149
      GO TO 806                                           MOM 150
804 CONTINUE                                           MOM 151
      GO TO 831                                           MOM 152
806 EMN(J)=WGHT(KK)                                     MOM 153
      NAMOLD=NAME(KK)                                    MOM 154
      WTOLD=WGHT(KK)                                    MOM 155
      40 EM(J)=EMN(J)/EN                                  MOM 156
      801 CONTINUE                                       MOM 157
C                                     MOM 158
C TRANSFER ACCORDING TO NTYPE, TO INPUT REMAINING DATA. MOM 159
C                                     MOM 160
      GO TO (100,200,300,400,500,600),NTYPE           MOM 161
C                                     MOM 162
C ERROR WRITE OUT, AND GO ON TO NEXT CASE.             MOM 163
C                                     MOM 164
      831 CONTINUE                                       MOM 165
      WRITE OUTPUT TAPE IT2,2016,MOL(J)                 MOM 166
      810 WRITE OUTPUT TAPE IT2,2020,(SUBS(I),I=1,4)    MOM 167
B      CHECK=230160606060                               MOM 168

```


TABLE III. - Continued. CODE LISTING

825	READ INPUT TAPE IT1,1003,(COMENT(J),J=1,14)	MOM	169
	DIFC=NCHECK-NCOM	MOM	170
B	IF(DIFC*7777777777777777)825,901,825	MOM	171
C		MOM	172
C	PLANAR SUBSTANCES INPUT	MOM	173
C		MOM	174
	100 READ INPUT TAPE IT1,1005,NCHAIN,NBRAN	MOM	175
	WRITE OUTPUT TAPE IT2,2100,NCHAIN,NBRAN,(COMENT(J),J=1,13)	MOM	176
	105 DO 101 J=1,NATMS	MOM	177
	101 Z(J)=0.	MOM	178
	X(1)=0.	MOM	179
	Y(1)=0.	MOM	180
	N=1	MOM	181
	IF(NCHAIN-2)132,102,102	MOM	182
	132 WRITE OUTPUT TAPE IT2,2018,NCHAIN	MOM	183
	GO TO 810	MOM	184
	102 NM1=NCHAIN-1	MOM	185
	NM2=NCHAIN-2	MOM	186
	READ INPUT TAPE IT1,1001,(BOND(J),J=1,NM1),(ANG(J),J=1,NM2)	MOM	187
	IF(ANG(1))115,116,115	MOM	188
	115 ANGLE=(180.0-ANG(1))*RAD	MOM	189
	116 X(2)=(BOND(1))*COSF(ANGLE)	MOM	190
	Y(2)=BOND(1)*SINF(ANGLE)	MOM	191
	ANGDEG=0.	MOM	192
	DO 112 J=3,NCHAIN	MOM	193
	ANGLE=ANGDEG*RAD	MOM	194
	SIGN=(-1)**(J-1)	MOM	195
	X(J)=X(J-1)+SIGN*BOND(J-1)*COSF(ANGLE)	MOM	196
	IF(MODF(ANGDEG,180.))114,113,114	MOM	197
	113 SIGN=0.	MOM	198
	114 Y(J)=Y(J-1)+SIGN*BOND(J-1)*SINF(ANGLE)	MOM	199
	112 ANGDEG=ANGDEG+ANG(J-1)	MOM	200
	WRITE OUTPUT TAPE IT2,2108,N,MOL(1),EMN(1),X(1),Y(1),Z(1),BOND(1)	MOM	201
	IF(NCHAIN-2)1044,1044,1045	MOM	202
	1045 WRITE OUTPUT TAPE IT2,2101,(J,MOL(J),ANG(J-1),EMN(J),X(J),Y(J),	MOM	203
	Z(J),BOND(J),J=2,NM1)	MOM	204
	1044 N=NCHAIN	MOM	205
	WRITE OUTPUT TAPE IT2,2102,N,MOL(N),EMN(N),X(N),Y(N),Z(N)	MOM	206
	N=N+1	MOM	207
	104 IF(NBRAN)103,135,103	MOM	208
	103 CONTINUE	MOM	209
	DO 130 L=1,NBRAN	MOM	210
	READ INPUT TAPE IT1,1005,NATCH,NBRATS	MOM	211
	IF(NCHAIN-NATCH)107,107,106	MOM	212
	107 WRITE OUTPUT TAPE IT2,2021,NCHAIN,NATCH	MOM	213
	GO TO 810	MOM	214
	106 READ INPUT TAPE IT1,1001,(BOND(J),J=1,NBRATS),(ANGBR(J),J=1,NBRATS)	MOM	215
	NP1=N+1	MOM	216
	NN=NATCH	MOM	217
	NNN=1	MOM	218
	M=N	MOM	219
	128 ASSIGN 127 TO NNNN	MOM	220
	IF(NATCH-2)121,122,123	MOM	221
	121 ANGDEG=ANGBR(1)-ANG(1)	MOM	222
	GO TO 125	MOM	223
	122 ANGDEG=ANGBR(1)	MOM	224

TABLE III. - Continued. CODE LISTING

GO TO 125	MOM 225
123 ANGDEG=ANGBR(1)	MOM 226
NAM1=NATCH-1	MOM 227
DO 124 J=2,NAM1	MOM 228
124 ANGDEG=ANGDEG+ANG(J)	MOM 229
125 SIGN=(-1.0)**NATCH	MOM 230
129 ANGLE=ANGDEG*RAD	MOM 231
X(M)=X(NN)+SIGN*BOND(NNN)*COSF(ANGLE)	MOM 232
Y(M)=Y(NN)+SIGN*BOND(NNN)*SINF(ANGLE)	MOM 233
IF(NBRATS-1)145,134,136	MOM 234
136 GO TO NNNN,(127,134)	MOM 235
127 ANGDEG=ANGDEG+ANGBR(2)	MOM 236
SIGN=-SIGN	MOM 237
NN=N	MOM 238
NNN=2	MOM 239
M=M+1	MOM 240
ASSIGN 134 TO NNNN	MOM 241
GO TO 129	MOM 242
134 WRITE OUTPUT TAPE IT2,2103,L,MOL(NATCH),NATCH	MOM 243
DO 131 KK=1,NBRATS	MOM 244
NN=N-1+KK	MOM 245
131 WRITE OUTPUT TAPE IT2,2104,NN,MOL(NN),BOND(KK),ANGBR(KK),EMN(NN),	MOM 246
IX(NN),Y(NN),Z(NN)	MOM 247
130 N=N+NBRATS	MOM 248
135 IF(SWITCH1401,800,401	MOM 249
145 WRITE OUTPUT TAPE IT2,2125,NBRATS,L,MOL(NATCH),NATCH,MOL(N),	MOM 250
IMOL(NP1)	MOM 251
GO TO 810	MOM 252
C	MOM 253
C INPUT FOR PYRAMIDAL XY3 ALWAYS 4 ATOMS-- X, Y, Y, Y	MOM 254
C BLN=X-Y BOND LENGTH, ALF= Y-X-Y ANGLE	MOM 255
C	MOM 256
200 CONTINUE	MOM 257
READ INPUT TAPE IT1,1001,BLN,ALF	MOM 258
WRITE OUTPUT TAPE IT2,2106,BLN,ALF,(COMENT(J),J=1,13)	MOM 259
ANGL=(ALF/2.)*RAD	MOM 260
SINAL=SINF(ANGL)	MOM 261
COSAL=COSF(ANGL)	MOM 262
SISQ=SINAL**2	MOM 263
COSQ=COSAL**2	MOM 264
X(1)=0.	MOM 265
X(2)=BLN*SINAL	MOM 266
X(3)=-X(2)	MOM 267
X(4)=0.	MOM 268
Y(1)=0.	MOM 269
Y(2)=0.	MOM 270
Y(3)=0.	MOM 271
TEM=(1.-3.*SISQ+COSQ)/(2.*COSAL)	MOM 272
Y(4)=SQRTF(BLN**2*(1.-TEM**2))	MOM 273
Z(1)=0.	MOM 274
Z(2)=BLN*COSAL	MOM 275
Z(3)=Z(2)	MOM 276
Z(4)=BLN*TEM	MOM 277
GO TO 815	MOM 278
C	MOM 279
C INPUT FOR TETRAHEDRAL-REGULAR XY4 ALWAYS 5 ATOMS---X Y Y Y Y	MOM 280

TABLE III. - Continued. CODE LISTING

C	BLN=X-Y BOND LENGTH	MOM	281
C		MOM	282
	300 CONTINUE	MOM	283
	READ INPUT TAPE IT1,1001,BLN	MOM	284
	WRITE OUTPUT TAPE IT2,2105,BLN,(COMENT(J),J=1,13)	MOM	285
	X(1)=0.	MOM	286
	X(2)=BLN*0.81649658	MOM	287
	X(3)=0.	MOM	288
	X(4)=-X(2)	MOM	289
	X(5)=0.	MOM	290
	Y(1)=BLN/3.	MOM	291
	Y(2)=0.	MOM	292
	Y(3)=0.	MOM	293
	Y(4)=0.	MOM	294
	Y(5)=BLN*1.33333333	MOM	295
	Z(1)=0.	MOM	296
	Z(2)=BLN*0.47140452	MOM	297
	Z(3)=-2.*Z(2)	MOM	298
	Z(4)=Z(2)	MOM	299
	Z(5)=0.	MOM	300
	GO TO 815	MOM	301
C		MOM	302
C	NON-PLANAR WITH OUT OF PLANE ATOMS	MOM	303
C		MOM	304
	400 CONTINUE	MOM	305
	READ INPUT TAPE IT1,1005,NCHAIN,NBRAN,NOPA	MOM	306
	WRITE OUTPUT TAPE IT2,2400,NCHAIN,NBRAN,NOPA,(COMENT(J),J=1,13)	MOM	307
	SWITCH=0.1	MOM	308
	GO TO 105	MOM	309
	401 DO 410 L=1,NOPA	MOM	310
	READ INPUT TAPE IT1,1007,NATCH,BLN,ANGDEG,SIGNOP	MOM	311
	ANGLE=ANGDEG*RAD	MOM	312
	GO TO (402,402,403),NATCH	MOM	313
	402 SIGN=+1.0	MOM	314
	GO TO 405	MOM	315
	403 SIGN=-1.0	MOM	316
	405 X(N)=X(NATCH)+SIGN*BLN*COSF(ANGLE)	MOM	317
	Y(N)=Y(NATCH)	MOM	318
	Z(N)=SIGNOP*BLN*SINF(ANGLE)	MOM	319
	404 WRITE OUTPUT TAPE IT2,2401,L,MOL(NATCH),NATCH	MOM	320
	WRITE OUTPUT TAPE IT2,2104,N,MOL(N),BLN,ANGDEG,EMN(N),X(N),Y(N),	MOM	321
	Z(N)	MOM	322
	410 N=N+1	MOM	323
	GO TO 800	MOM	324
C		MOM	325
C	COORDINATES READ IN--U+V*COS(ALPHA)	MOM	326
C		MOM	327
	500 CONTINUE	MOM	328
	WRITE OUTPUT TAPE IT2,2500,(COMENT(J),J=1,13)	MOM	329
	DO 501 L=1,NATMS	MOM	330
	READ INPUT TAPE IT1,1500,(BOND(J),V(J),ANG(J),J=1,3)	MOM	331
	WRITE OUTPUT TAPE IT2,2501,MOL(L),(BOND(J),V(J),ANG(J),J=1,3)	MOM	332
	X(L)=BOND(1)+V(1)*(COSF(ANG(1)*RAD))	MOM	333
	Y(L)=BOND(2)+V(2)*(COSF(ANG(2)*RAD))	MOM	334
	Z(L)=BOND(3)+V(3)*(COSF(ANG(3)*RAD))	MOM	335
	501 CONTINUE	MOM	336

TABLE III. - Continued. CODE LISTING

GO TO 815	MOM 337
C	MOM 338
C COORDINATES READ IN---X, Y, Z	MOM 339
C SPECIAL FOR MOLECULES NOT CATEGORIZED	MOM 340
C	MOM 341
600 CONTINUE	MOM 342
READ INPUT TAPE IT1,1001,(X(J),J=1,NATMS)	MOM 343
READ INPUT TAPE IT1,1001,(Y(J),J=1,NATMS)	MOM 344
READ INPUT TAPE IT1,1001,(Z(J),J=1,NATMS)	MOM 345
WRITE OUTPUT TAPE IT2,2600,(COMENT(J),J=1,13)	MOM 346
815 WRITE OUTPUT TAPE IT2,2019,(J,MOL(J),EMN(J),X(J),Y(J),Z(J),	MOM 347
1J=1,NATMS)	MOM 348
C	MOM 349
C COMPUTE ELEMENTS OF SYMMETRIC MATRIX, A.	MOM 350
C	MOM 351
800 CONTINUE	MOM 352
EMM=0.	MOM 353
SMX=0.	MOM 354
SMY=0.	MOM 355
SMZ=0.	MOM 356
SXY=0.	MOM 357
SXZ=0.	MOM 358
SYZ=0.	MOM 359
SMXY=0.	MOM 360
SMXZ=0.	MOM 361
SMYZ=0.	MOM 362
DO 10 J=1,NATMS	MOM 363
EMM=EMM+EM(J)	MOM 364
EMX=EM(J)*X(J)	MOM 365
EMY=EM(J)*Y(J)	MOM 366
XSQ=X(J)**2	MOM 367
YSQ=Y(J)**2	MOM 368
ZSQ=Z(J)**2	MOM 369
SMX=SMX+EMX	MOM 370
SMY=SMY+EMY	MOM 371
SMZ=SMZ+(EM(J)*Z(J))	MOM 372
SMXY=SMXY+(EMX*Y(J))	MOM 373
SMXZ=SMXZ+(EMX*Z(J))	MOM 374
SMYZ=SMYZ+(EMY*Z(J))	MOM 375
SXY=SXY+(EM(J)*(XSQ+YSQ))	MOM 376
SXZ=SXZ+(EM(J)*(XSQ+ZSQ))	MOM 377
SYZ=SYZ+(EM(J)*(YSQ+ZSQ))	MOM 378
10 CONTINUE	MOM 379
SMXSQ=SMX**2	MOM 380
SMYSQ=SMY**2	MOM 381
SMZSQ=SMZ**2	MOM 382
EY11=SYZ-((SMYSQ+SMZSQ)/EMM)	MOM 383
EY22= SXZ-((SMXSQ+SMZSQ)/EMM)	MOM 384
EY33= SXY-((SMYSQ+SMXSQ)/EMM)	MOM 385
EY12=SMX*SMY/EMM-SMXY	MOM 386
EY13=SMX*SMZ/EMM-SMXZ	MOM 387
EY23=SMY*SMZ/EMM-SMYZ	MOM 388
A(1,1)=EY11	MOM 389
A(2,2)=EY22	MOM 390
A(3,3)=EY33	MOM 391
A(1,2)=EY12	MOM 392

TABLE III. - Continued. CODE LISTING

```

      A(2,1)=EY12                                MOM 393
      A(1,3)=EY13                                MOM 394
      A(3,1)=EY13                                MOM 395
      A(2,3)=EY23                                MOM 396
11  A(3,2)=EY23                                MOM 397
C
C GET EIGENVALUES.                                MOM 398
C                                                MOM 399
C                                                MOM 400
      CALL JACOBI(3,1.0E-9)                      MOM 401
      PRODD=D(1,1)*D(2,2)*D(3,3)                 MOM 402
C                                                MOM 403
C WRITE OUT MATRICES AND EIGENVALUES.             MOM 404
C                                                MOM 405
      WRITE OUTPUT TAPE IT2,2200,EY11,EY12,EY13,D(1,1),D(2,2),EY12,EY22,MOM 406
      1EY23,D(3,3),EY13,EY23,EY33,PRODD          MOM 407
820  WRITE OUTPUT TAPE IT2,2201,(((D(I,J),J=1,3),(S(I,J),J=1,3)),I=1,3)MOM 408
      WRITE OUTPUT TAPE IT2,2202,((SD(I,J),J=1,3),I=1,3)MOM 409
C                                                MOM 410
C GO TO READ DATA FOR A NEW CASE.                MOM 411
C                                                MOM 412
      GO TO 900                                    MOM 413
1000 FORMAT(2I3,4A6,25A2)                        MOM 414
1001 FORMAT(8F10.4)                              MOM 415
1002 FORMAT(13,9F8.3/(10F8.3))                   MOM 416
1003 FORMAT(13A6,A2)                              MOM 417
1005 FORMAT(3I3)                                  MOM 418
1006 FORMAT(40A2)                                  MOM 419
1007 FORMAT(13,7X,3F10.4)                         MOM 420
1500 FORMAT(9F8.4)                                MOM 421
2015 FORMAT(1H1,15X,33HPRINCIPAL MOMENTS OF INERTIA OF 4A6)MOM 422
2016 FORMAT(13HL THIS ATOM A6,36HIS NOT IN THE NAME AND WEIGHT TABLE.)MOM 423
2018 FORMAT(52HL NO. OF CHAIN ATOMS IS LESS THAN 2. NCHAIN= 15)MOM 424
2019 FORMAT(1HK,5X,3HNO.,6X,4HATOM,7X,16HMOLECULAR WEIGHT,6X,1HX,16X, MOM 425
      11HY,16X,1HZ/(1HJ,17,8X,A6,4X,4G17.8))    MOM 426
2020 FORMAT(30HL THIS CASE, SUBSTANCE IS 4A6,48H, CANNOT BE CONTINUUMOM 427
      1ED, GO ON TO NEXT SUBSTANCE. )            MOM 428
2021 FORMAT(58HL BRANCH IS ATTACHED TO LAST OR NO CHAIN ATOM. THERE AMOM 429
      1RE 13,54H CHAIN ATOMS, THE BRANCH IS ATTACHED TO CHAIN ATOM NO.13)MOM 430
2100 FORMAT(14HK PLANAR WITH 13,16H CHAIN ATOMS AND 13,10H BRANCHES., MOM 431
      14X, 13A6)                                  MOM 432
2101 FORMAT(18,8X,A6,15X,F9.2,4X,4G17.8/F35.4)    MOM 433
2102 FORMAT(18,8X,A6,28X,4G17.8)                  MOM 434
2103 FORMAT(1HK,6X,10HBRANCH NO. 13,26H, ATTACHED TO CHAIN ATOM A6, MOM 435
      1 9H, NUMBER 13/1HJ,5X,3HNO.,6X,4HATOM,7X,11HBOND LENGTH,4X, MOM 436
      25HANGLE /)                                  MOM 437
2104 FORMAT(18,8X,A6,F13.4,F11.2,4X,4G17.8)       MOM 438
2105 FORMAT(40HL TETRAHEDRAL-REGULAR WITH BOND LENGTH= F8.4 ,3X, MOM 439
      113A6)                                       MOM 440
2106 FORMAT(30HL PYRAMIDAL WITH BOND LENGTH= F10.5,13H , AND ANGLE= MOM 441
      1F10.2/40X, 13A6,A2)                       MOM 442
2107 FORMAT(1HJ,40X,13A6,A2)                     MOM 443
2108 FORMAT(1HK,5X,60HNO. CHAIN ATOM BOND LENGTH ANGLE MOLECMOM 444
      1ULAR WEIGHT,7X,1HX,16X,1HY,16X,1HZ/1HJ,17,8X,A6,28X,4G17.8/F35.4) MOM 445
2125 FORMAT(61HL NO. OF BRANCH ATOMS IS =, OR LESS THAN 0. NBRANCH ATMOM 446
      1OMS= 14,20H THIS IS BRANCH NO. 14/26H, ATTACHED TO CHAIN ATOM 6, MOM 447
      29H, NUMBER 14,30H. THE ATOMS ON THIS BRANCH ARE 2A6) MOM 448

```


TABLE III. - Continued. CODE LISTING

```

2200 FORMAT(44HL THE GENERATED SYMMETRIC MATRIX IS .....36X,      MOM 449
2 46HTHE EIGENVALUES, LAMBDA, OF THE MATRIX ARE THE/80X,      MOM 450
3 39HPRINCIPAL MOMENTS OF INERTIA (G-SQ CM)/8X,1HI,55X,1HI/      MOM 451
4 8X,1HI,3G17.7,4X,1HI/8X,1HI,55X,1HI,18X,19HLAMBDA(1)*1.0E+39 =      MOM 452
5 G16.8/8X,1HI,55X,1HI,18X,19HLAMBDA(2)*1.0E+39 =G16.8/8X,1HI,      MOM 453
6 3G17.7,4X,1HI,18X,19HLAMBDA(3)*1.0E+39 =G16.8/8X,1HI,55X,1HI/      MOM 454
7 8X,1HI,55X,1HI/8X,1HI,3G17.7,4X,1HI,18X,26HPRODUCT LAMBDA*1.0E+1      MOM 455
817 =G16.8/8X,1HI,55X,1HI)      MOM 456
2201 FORMAT(1HI,4X,61HTHE GENERATED MATRICES FROM THE JACOBI METHOD ARE      MOM 457
1 AS FOLLOWS,/1HK,4X, 24HTHE DIAGONAL MATRIX (D) ,40X, 27HTHE EIGEN      MOM 458
2VECTOR MATRIX (S) / 2HJI,58X,1HI, 9X,1HI,58X,1HI/      MOM 459
3 2H I,G16.8,2G20.8,2X,1HI, 9X,1HI,G16.8,2G20.8,2X,1HI/      MOM 460
4 2H I,58X,1HI,9X,1HI,58X,1HI/      MOM 461
5 2H I,G16.8,2G20.8,2X,1HI, 9X,1HI,G16.8,2G20.8,2X,1HI/      MOM 462
6 2H I,58X,1HI,9X,1HI,58X,1HI/      MOM 463
7 2H I,G16.8,2G20.8,2X,1HI, 9X,1HI,G16.8,2G20.8,2X,1HI/      MOM 464
8 2H I,58X,1HI,9X,1HI,58X,1HI)      MOM 465
2202 FORMAT(1HL,4X, 39HTHE MATRIX (S)*(D)*(S TRANSPOSE), WHICH /      MOM 466
19X,36HSHOULD EQUAL THE ORIGINAL MATRIX IS,/      MOM 467
22HJI,58X,1HI/      MOM 468
32H I,G16.8,2G20.8,2X,1HI/      MOM 469
42H I,58X,1HI/      MOM 470
52H I,G16.8,2G20.8,2X,1HI/      MOM 471
62H I,58X,1HI/      MOM 472
72H I,G16.8,2G20.8,2X,1HI/      MOM 473
82H I,58X,1HI)      MOM 474
2400 FORMAT(1HL,10X,15HNON-PLANAR WITH I3,14H CHAIN ATOMS, I3,      MOM 475
122H PLANAR BRANCHES, AND I3,21H OUT-OF-PLANE ATOMS. /1HJ,40X,      MOM 476
213A6)      MOM 477
2401 FORMAT(1HK,6X,17H O-P-A BRANCH NO.I3,26H, ATTACHED TO CHAIN ATOM      MOM 478
1A6, 8H, NUMBER I3/1HJ,5X,3HNO.,6X,4HATOM,7X,11HBOND LENGTH,4X,11HOMOM      MOM 479
2-P-A ANGLE /)      MOM 480
2500 FORMAT(1HL, 5X,35HCOORDINATES READ IN-- U+VCDS(ALPHA),5X,13A6/      MOM 481
11HL,19HU,V,ALF INPUT X,45X,1HY,45X,1HZ/6HJ ATOM,6X,1HU,10X,1HV      MOM 482
2,10X,5HALPHA,16X,1HU,10X,1HV,10X,5HALPHA,16X,1HU,10X,1HV,10X,      MOM 483
35HALPHA / )      MOM 484
2501 FORMAT(2X,A6,3F11.4,10X,3F11.4,10X,3F11.4)      MOM 485
2600 FORMAT(1HL, 5X,35HCOORDINATES READ IN-- X, Y, Z 13A6)      MOM 486
END(0,1,0,1,0,1,1,0,0,1,0,0,0,0,0)

```


TABLE III. - Continued. CODE LISTING

C	SUBROUTINE JACOBI(NN,FACTOR)	JAC 001
C	SUBROUTINE JACOBI, COMPUTES ALL EIGENVALUES AND THE EIGENVECTOR	JAC 002
C	MATRIX OF REAL SYMMETRIC MATRICES	JAC 003
C	CALLING PROGRAM MUST,	JAC 004
C	1 HAVE COMMON Y,S,SD	JAC 005
C	2 HAVE DIMENSION Y(3,3),S(3,3),SD(3,3)	JAC 006
C	ORIGINAL MATRIX A(I,J) IN ARRAY Y.	JAC 007
C		JAC 008
C	3 CALLING SEQUENCE ,	JAC 009
C	CALL JACOBI(N,FACTOR),WHERE N= ORDER OF MATRIX	JAC 010
C	FACTOR= MAGNITUDE OF FINAL THRESHOLD	JAC 011
C	N IS AN INTEGER, FACTOR IS A FLOATING POINT NO.	JAC 012
C		JAC 013
C	RETURN IS MADE TO CALLING PROGRAM WITH,	JAC 014
C	1 DIAGONAL MATRIX (D) IN ARRAY Y, THE DIAGONAL ELEMENTS ARE THE	JAC 015
C	EIGENVALUES.	JAC 016
C	THE ORIGINAL MATRIX, (A(I,J)), IS DESTROYED IN THE	JAC 017
C	COMPUTATION PROCESS.	JAC 018
C		JAC 019
C	2 EIGENVECTOR MATRIX (S) IN ARRAY S.	JAC 020
C		JAC 021
C	3 MATRIX (S=D*S TRANSPOSE), WHICH SHOULD EQUAL THE ORIGINAL	JAC 022
C	MATRIX, IN ARRAY SD.	JAC 023
C		JAC 024
C	COMMON Y,S,SD	JAC 025
C	EQUIVALENCE (Y,D)	JAC 026
C	DIMENSION Y(3,3),D(3,3),S(3,3),SD(3,3) ,SDPR(3,3)	JAC 027
C	1 N=NN	JAC 028
C	2 SIGMA=N	JAC 029
C	DO 200 K=1,N	JAC 030
C	DO 200 L=1,N	JAC 031
C	SD(K,L)=0.	JAC 032
C	SDPR(K,L)=0.	JAC 033
C	200 S(K,L)=0.	JAC 034
C	DO 201 K=1,N	JAC 035
C	201 S(K,K)=1.0	JAC 036
C	3 IND=0	JAC 037
C	NM1=N-1	JAC 038
C	K=2	JAC 039
C	SUM=0.	JAC 040
C	DO 300 I=1,NM1	JAC 041
C	DO 301 J=K,N	JAC 042
C	301 SUM=SUM+Y(I,J)**2	JAC 043
C	K=K+1	JAC 044
C	300 CONTINUE	JAC 045
C	4 VI=SQRTF(2.*SUM)	JAC 046
C	IF(VI)5,80,5	JAC 047
C	5 V=VI	JAC 048
C	VF=VI*FACTOR	JAC 049
C	6 V=V/SIGMA	JAC 050
C	7 J=2	JAC 051
C	8 I=1	JAC 052
C	9 IF(ABSF(Y(I,J))-V)22,10,10	JAC 053
C	10 IND=1	JAC 054
C	11 ALAM=-Y(I,J)	JAC 055
C	12 AMU=(Y(I,1)-Y(J,J))/2.	JAC 056

TABLE III. - Continued. CODE LISTING

IF(AMU)111,112,112	JAC 057
111 SGNMU=-1.	JAC 058
GO TO 113	JAC 059
112 SGNMU=+1.	JAC 060
113 OMEGA=SGNMU*ALAM/(SQRTF(ALAM**2+AMU**2))	JAC 061
SINTH=OMEGA/(SQRTF(2.*(1.+SQRTF(1.-OMEGA**2))))	JAC 062
SISQ=SINTH*SINTH	JAC 063
COSTH=SQRTF(1.-SISQ)	JAC 064
COSQ=COSTH*COSTH	JAC 065
SICO=SINTH*COSTH	JAC 066
II=1	JAC 067
13 IF(II-I)130,14,130	JAC 068
130 IF(II-J)131,14,131	JAC 069
131 B1P=Y(II,I)*COSTH-Y(II,J)*SINTH	JAC 070
B1Q=Y(II,I)*SINTH+Y(II,J)*COSTH	JAC 071
Y(II,I)=B1P	JAC 072
Y(II,J)=B1Q	JAC 073
14 SMIP=S(II,I)*COSTH-S(II,J)*SINTH	JAC 074
SMIO=S(II,I)*SINTH+S(II,J)*COSTH	JAC 075
S(II,I)=SMIP	JAC 076
S(II,J)=SMIO	JAC 077
15 IF(N-II)17,17,16	JAC 078
16 II=II+1	JAC 079
GO TO 13	JAC 080
17 BPP=Y(I,I)*COSQ+Y(J,J)*SISQ-2.*Y(I,J)*SICO	JAC 081
BQQ=Y(I,I)*SISQ+Y(J,J)*COSQ+2.*Y(I,J)*SICO	JAC 082
BPQ=(Y(I,I)-Y(J,J))*SICO+Y(I,J)*(COSQ-SISQ)	JAC 083
Y(I,I)=BPP	JAC 084
Y(J,J)=BQQ	JAC 085
Y(I,J)=BPQ	JAC 086
18 II=1	JAC 087
19 Y(I,II)=Y(II,I)	JAC 088
Y(J,II)=Y(II,J)	JAC 089
20 IF(N-II)22,22,21	JAC 090
21 II=II+1	JAC 091
GO TO 19	JAC 092
22 IF(I-(J-1))23,24,24	JAC 093
23 I=I+1	JAC 094
GO TO 9	JAC 095
24 IF(N-J)26,26,25	JAC 096
25 J=J+1	JAC 097
GO TO 8	JAC 098
26 IF(IND-1)28,27,28	JAC 099
27 IND=0	JAC 100
GO TO 7	JAC 101
28 IF(V-VF)29,29,6	JAC 102
C IF OFF DIAGONAL ELEMENTS OF (D) MATRIX ARE LESS THAN 1.0E-19, SET =0.	JAC 103
29 CONTINUE	JAC 104
IF(ABSF(D(1,2))-1.0E-19)950,951,951	JAC 105
950 D(1,2)=0.	JAC 106
D(2,1)=0.	JAC 107
951 IF(ABSF(D(1,3))-1.0E-19)952,953,953	JAC 108
952 D(1,3)=0.	JAC 109
D(3,1)=0.	JAC 110
953 IF(ABSF(D(2,3))-1.0E-19)954,955,955	JAC 111
954 D(2,3)=0.	JAC 112

TABLE III. - Continued. CODE LISTING

D(3,2)=0.	JAC 113
955 CONTINUE	JAC 114
DO 50 I=1,N	JAC 115
DO 50 J=1,N	JAC 116
DO 50 K=1,N	JAC 117
50 SDPR(I,J)=SDPR(I,J)+S(I,K)*D(K,J)	JAC 118
DO 70 I=1,N	JAC 119
DO 70 J=1,N	JAC 120
DO 70 K=1,N	JAC 121
70 SD(I,J)=SD(I,J)+SDPR(I,K)*S(J,K)	JAC 122
71 RETURN	JAC 123
80 DO 81 I=1,N	JAC 124
81 SD(I,I)=Y(I,I)	JAC 125
GO TO 71	JAC 126
END(0,1,0,1,0,1,0,0,0,1,0,0,0,0,0)	

TABLE III. - Concluded. CODE LISTING

```

*DATA
102 6.02322 1.008 4.003 6.94 9.013 10.82 12.011 14.008 16.0
19.0 20.183 22.991 24.32 26.98 28.09 30.975 32.066 35.457 39.944
39.1 40.08 44.96 47.9 50.95 52.01 54.94 55.85 58.94 58.71
63.54 65.38 69.72 72.6 74.92 78.96 79.916 83.8 85.48 87.63
88.91 91.22 92.91 95.95 99.0 101.1 102.91 106.4 107.88 112.41
114.82 118.7 121.76 127.61 126.91 131.3 132.91 137.36 138.92 140.13
140.91 144.27 147.0 150.35 152.0 157.26 158.93 162.51 164.94 167.27
168.94 173.04 174.99 178.5 180.95 183.86 186.22 190.2 192.2 195.09
197.0 200.61 204.39 207.21 208.99 210.0 210.0 222.0 223.0 226.0
227.0 232.0 231.0 238.0 237.0 242.0 243.0 247.0 249.0 251.0
254.0 253.0 256.0 254.0
H HELIBEB C N O F NENAMGALSIP S CLARK CASCTIV CRMNFECONICUZNGAGEASSEBRKRRBSRY ZR
NBMTICRURHPDAGCDINSNSBTEI XECSBALACEPRNDPMSMEUGDTB0YHOERTMYBLUHFTAW REOSIRPTAUHG
TLPB8IPUATKNFRRAACTHPAU NPPUAMCMBKCFESFMMND0
WHITE JCP V32 P488 FEB 1960. C1
4 1HB02 H O B O
4 0
1.0 1.34 1.20 120. 180.
WHITE JCP V32 P488 FEB. 1960 B-O 1.36A, O-H 1.A, CYCLIC C1
12 1 (HB02)3 B O B O B O O H O H
6 3
1.36 1.36 1.36 1.36 1.36 120. 120. 120.
120.
1 2
1.36 1.0 120. 240.
3 2
1.36 1.0 120. 240.
5 2
1.36 1.0 120. 240.
KISLIUK JCP V18 P1109 AUG 1950. C1
4 2PCL3 P CLCLCL
2.043 100.1
BARTELL, JCP V23 P1854 OCT 1955. C1
5 3CCL4 C CLCLCLCL
1.760
MAXWELL JCP V3 P699 NOV 1935. C1
5 3P4 P P P P
1.353343
ESTIMATES FROM NAUH AND LIOH, (LIOH)2. C1
6 4(NAOH)2 NAO O NAH H
3 1 2
2.25 3.4472 40.0
2 1
2.25 40.0
2 0.96 120.0 +1.0
3 0.96 120.0 -1.0
BOWEN, CHEM. SOC., SPECIAL PUBL. NO. 11, 1958. C1
4 4S2CL2 CLS S CL
3 0 1
1.99 2.05 104.
3 1.99 104.0 -1.0
DODD TRANS FAR SOC V52 P1052 1956. FICTITIOUS THIRD ATOM. C1
6 4SF4 F S F F F
3 1 2
1.58 .79 60.
2 1
1.58 60.
2 1.58 110. 1.
2 1.58 110. -1.
PALMER JALS V60 P2360 1938. S-O 1.45A, S-CL 2.07A, O-S-CL 106, CL-S-CL 114. C1
4 5SOCL2 S O CLCL
0.
-7.73836 1.250594
2.07 57. 2.07 33.
2.07 57. -2.07 33.
BOWEN, CHEM. SOC., SPECIAL PUBL. NO. 11, 1958. C1
4 6S2CL2 S S CLCL
0.0 2.05 -.481424572.5314208
0.0 0.0 1.930888570.0
0.0 0.0 0.0 1.9308885

```

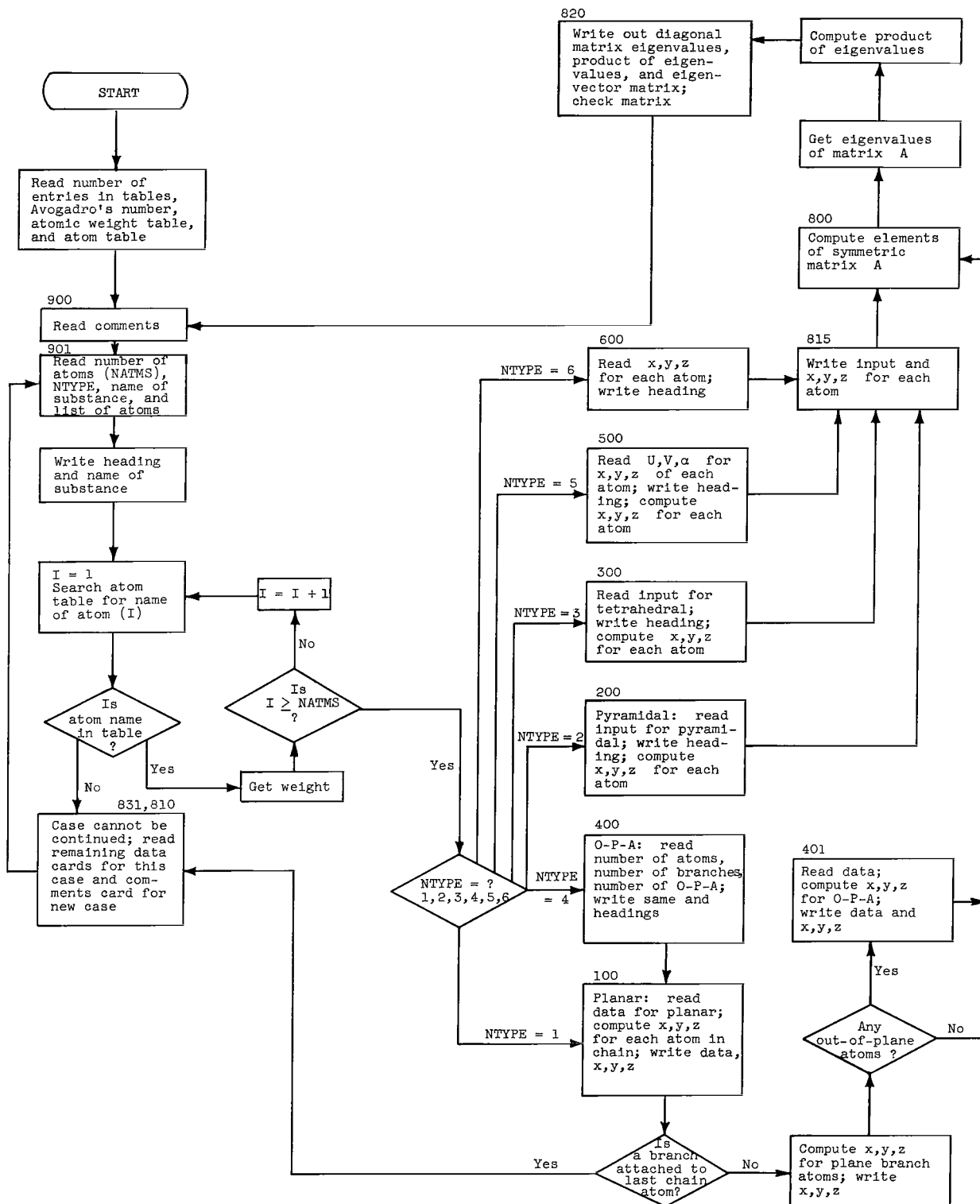
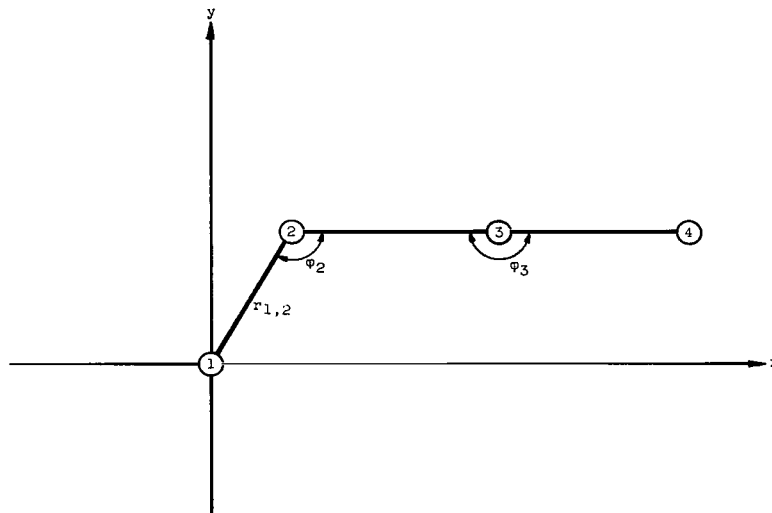
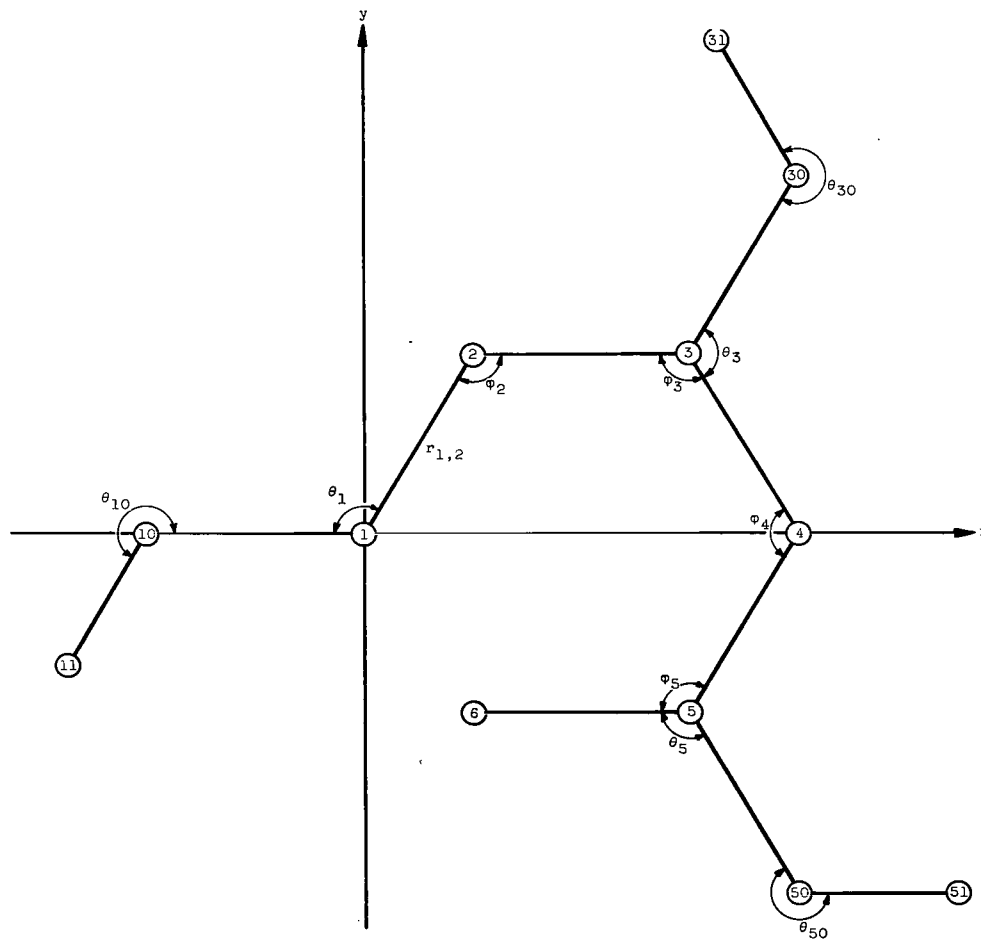



Figure 1. - Flow diagram of main program.



(a-1) No branching.

CD-7720

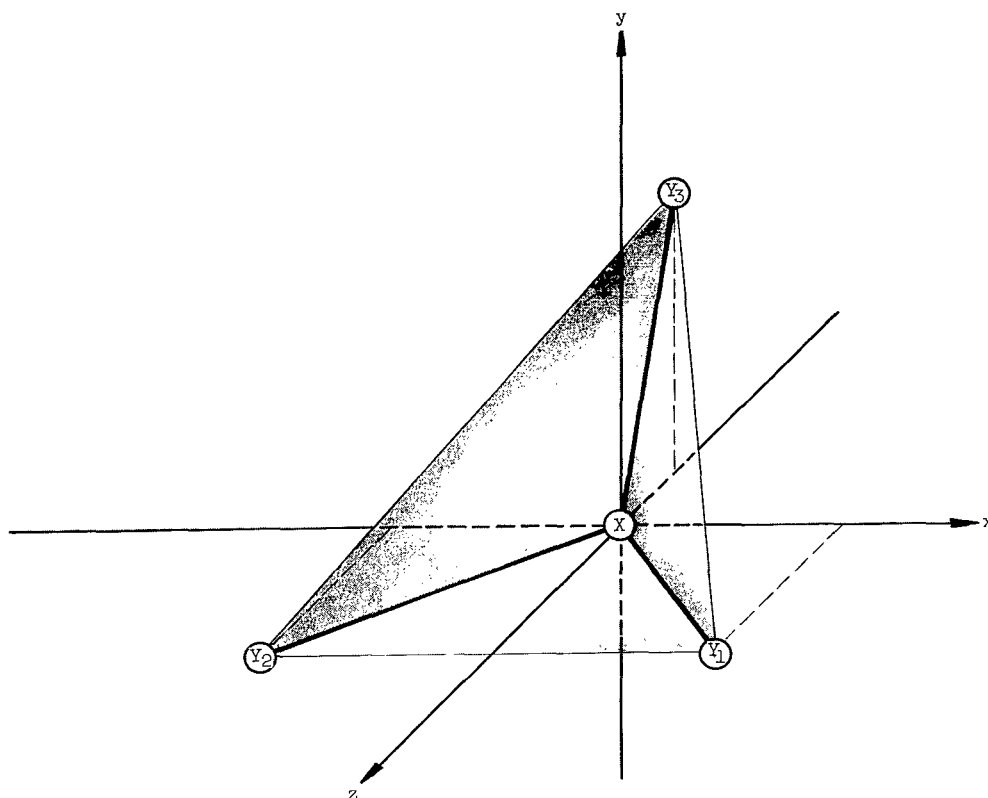


(a-2) With branching.

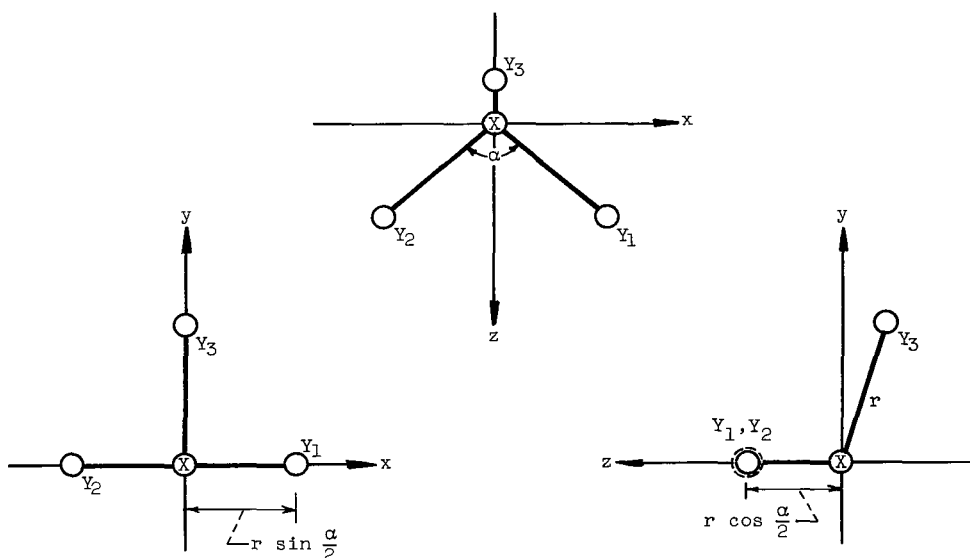
(a) Type 1 - planar.

Figure 2. - Diagrams of geometric types.

CD-7721



(b-1) Three-dimensional view.

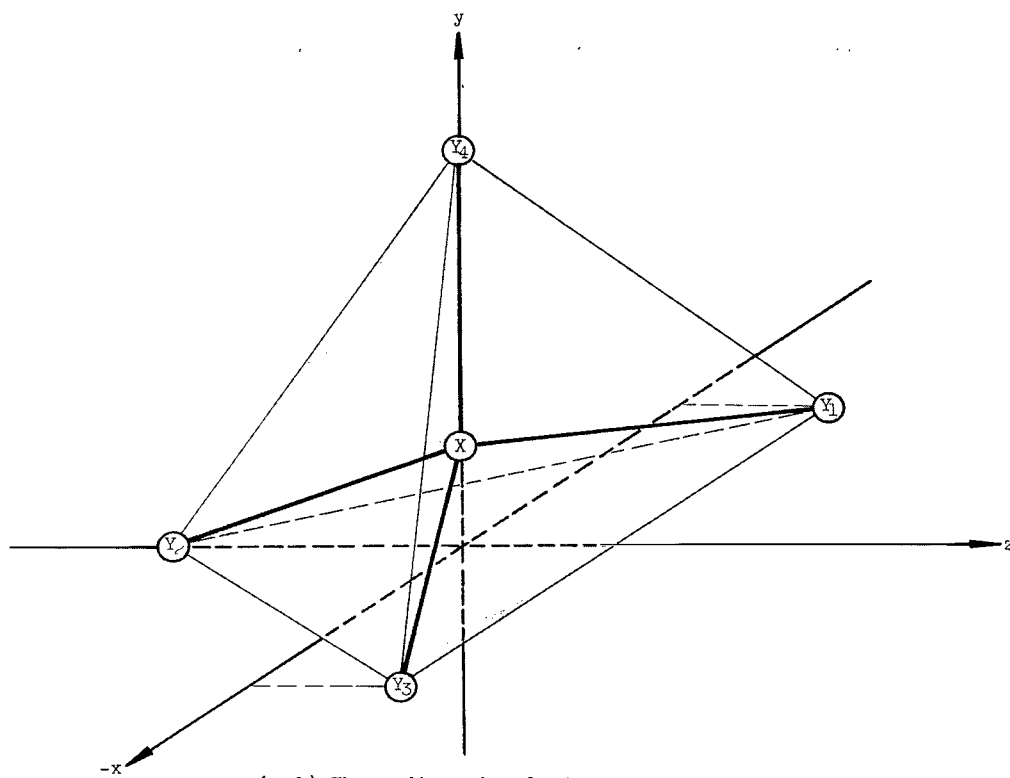


(b-2) Two-dimensional views.

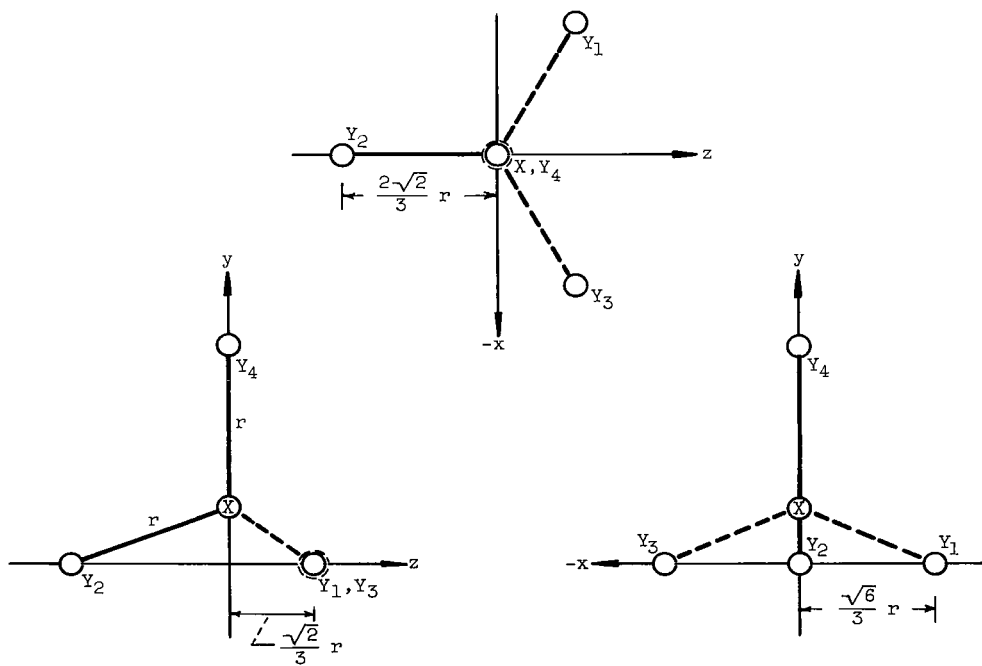
(b) Type 2 - pyramidal XY_3 .

Figure 2. - Continued. Diagrams of geometric types.

CD-7722



(c-1) Three-dimensional view.

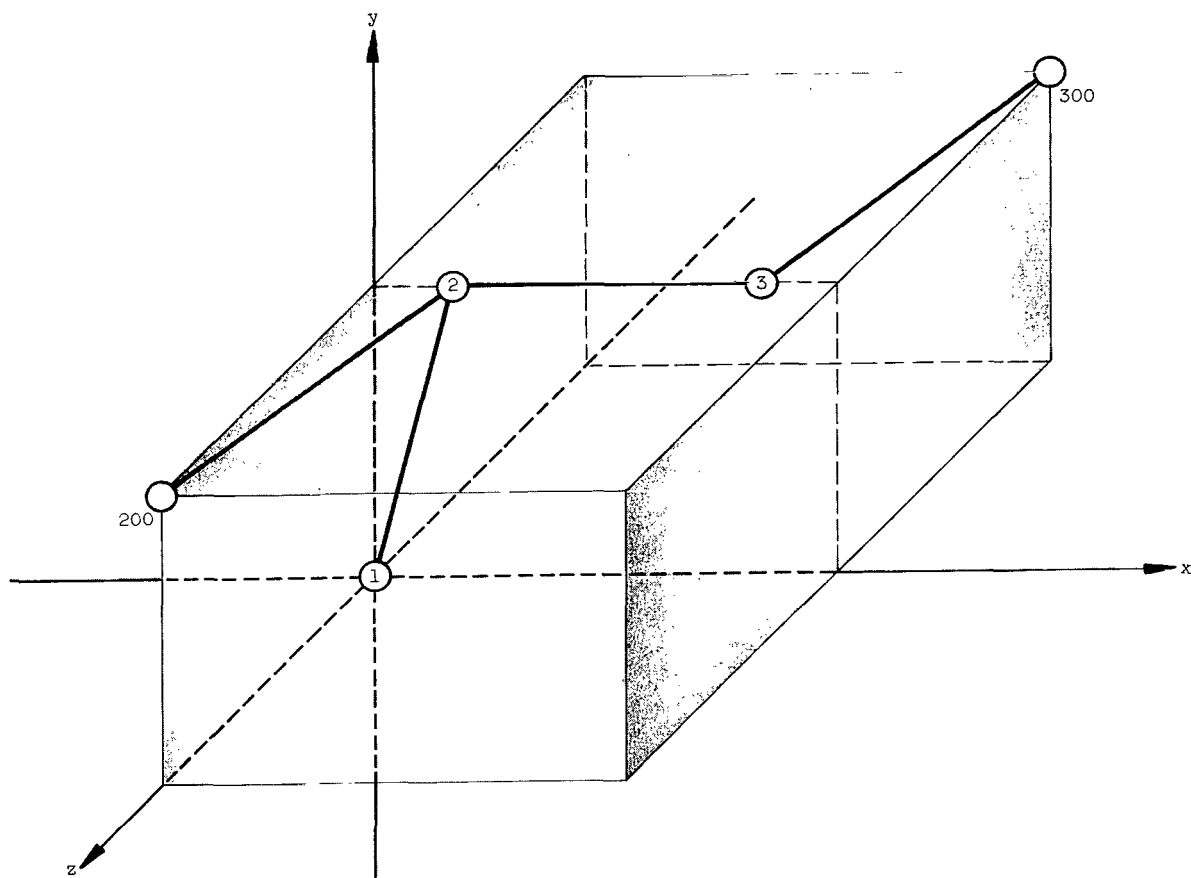


(c-2) Two-dimensional view.

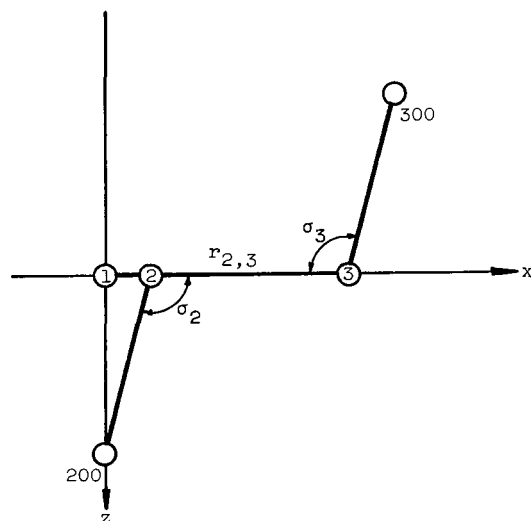
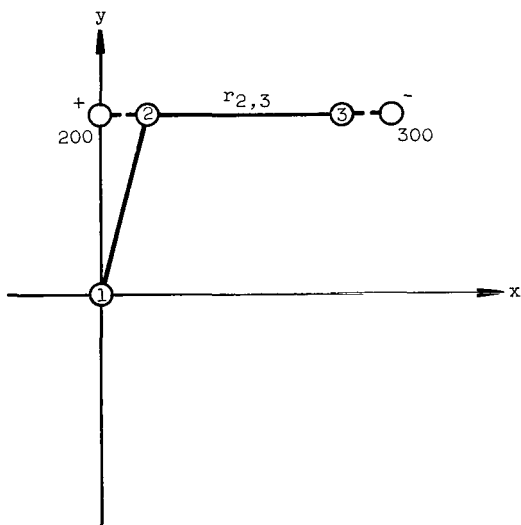
(c) Type 3 - tetrahedral XY_4 .

Figure 2. - Continued. Diagrams of geometric types.

CD-7724



(d-1) Three-dimensional view.

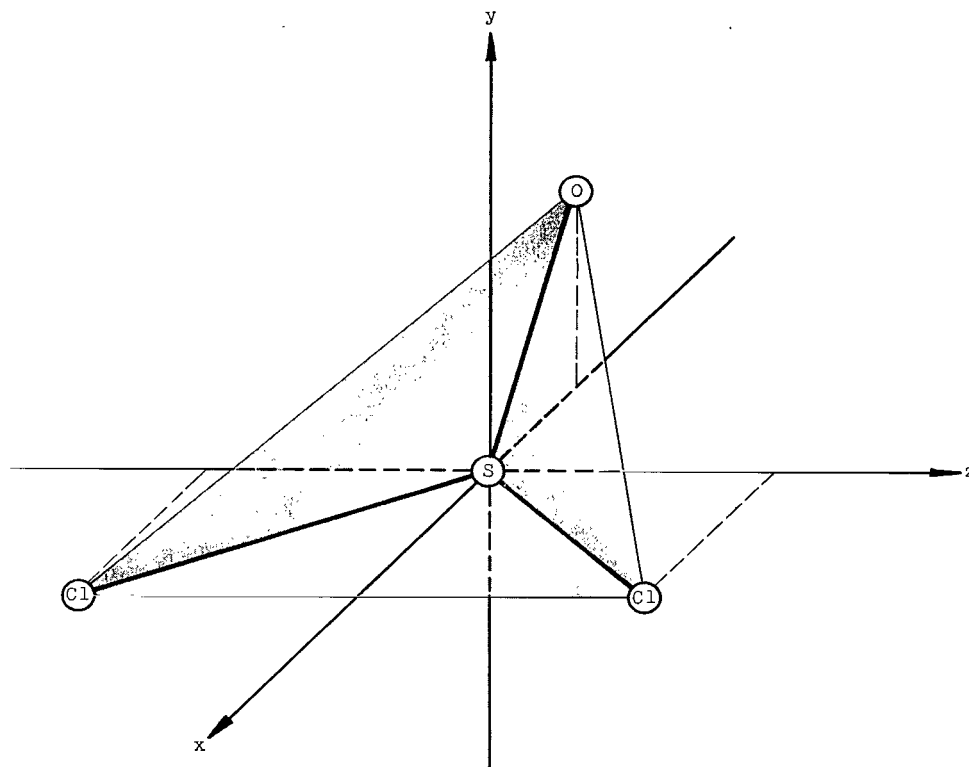


(d-2) Two-dimensional views.

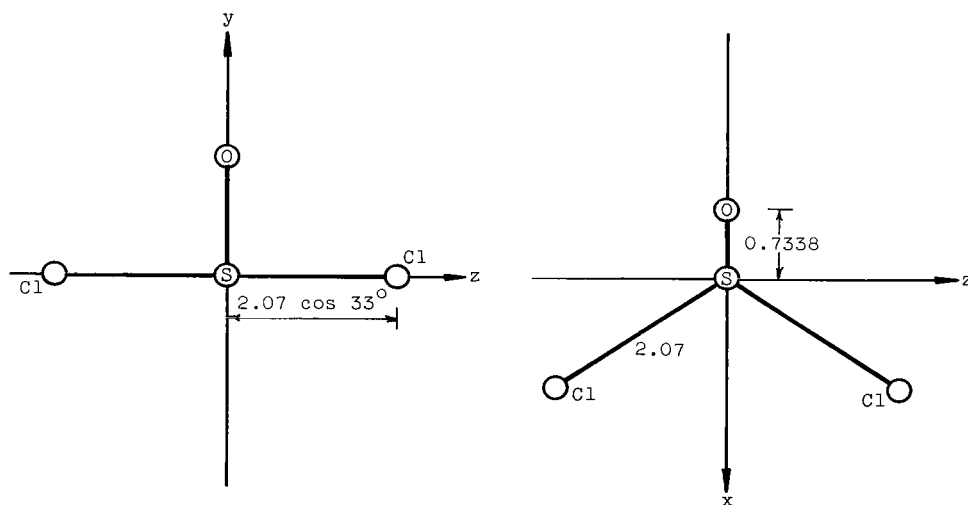
(d) Type 4 - other nonplanar.

Figure 2. - Continued. Diagrams of geometric types.

CD-7723



(e-1) Three-dimensional view.

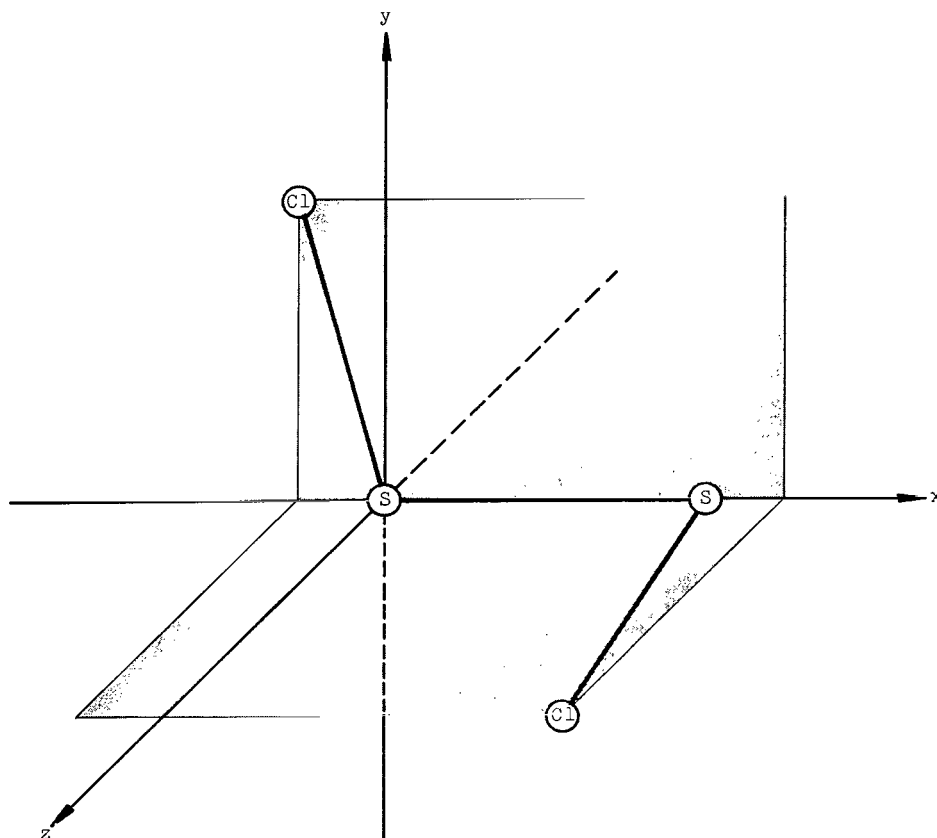


(e-2) Two-dimensional views.

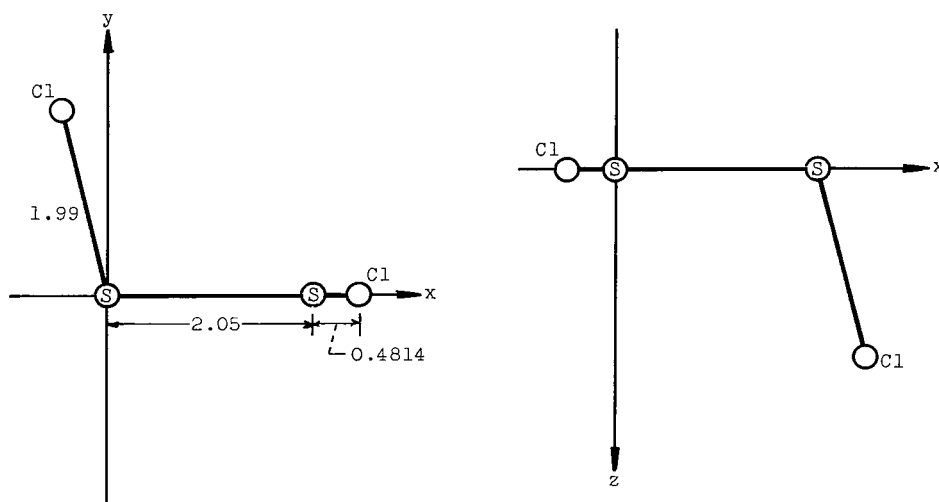
(e) Type 5 - general: $U + V \cos \alpha$; SOCl_2 .

Figure 2. - Continued. Diagrams of geometric types.

CD-7725



(f-1) Three-dimensional view.

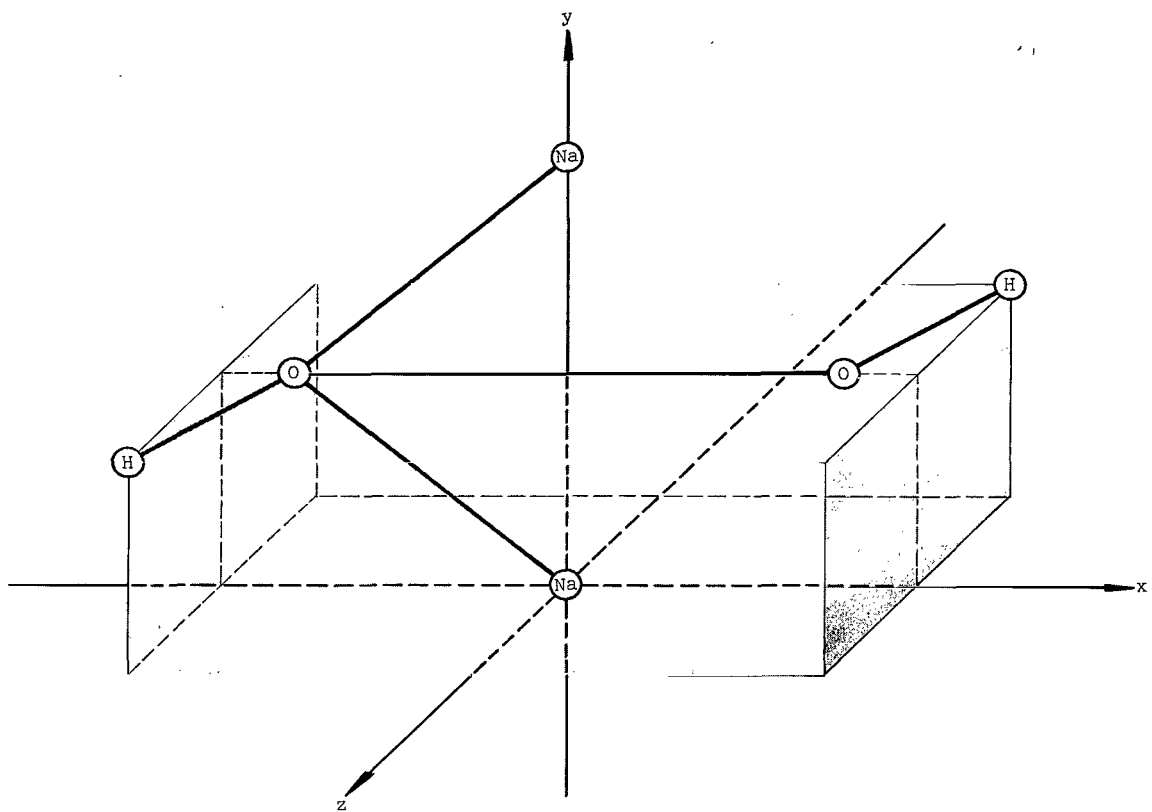


(f-2) Two-dimensional views.

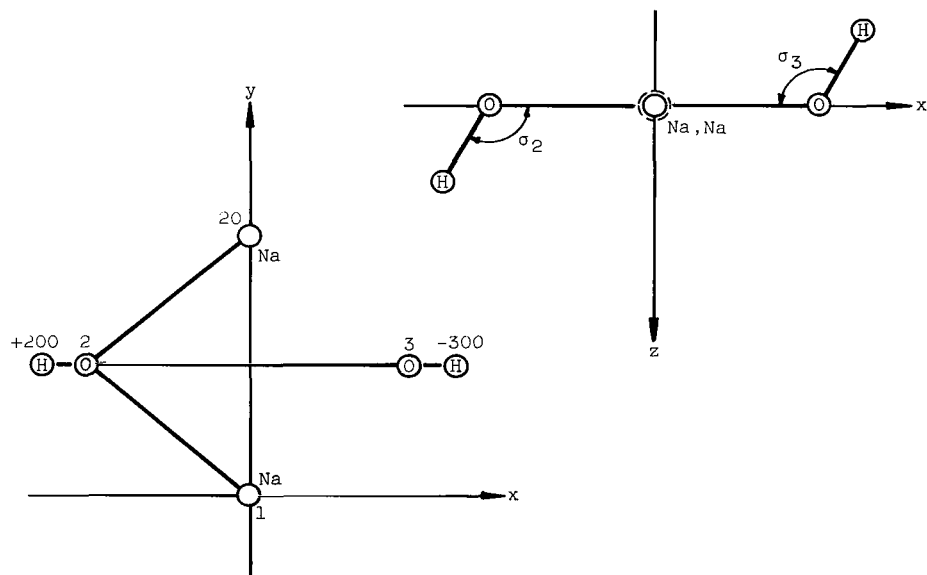
(f) Type 6 - general: $x, y, z; S_2Cl_2$.

Figure 2. - Concluded. Diagrams of geometric types.

CD-7726



(a-1) Three-dimensional view.

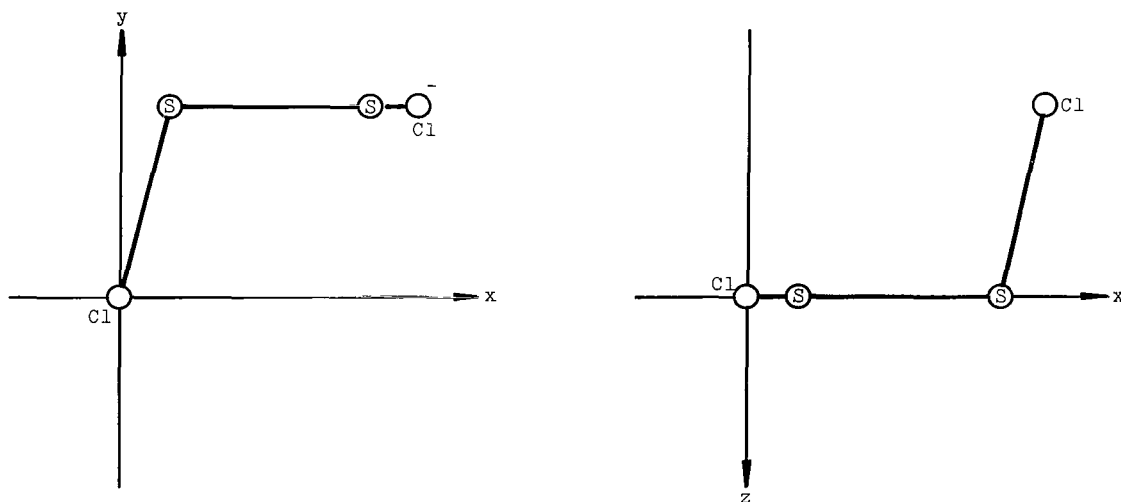
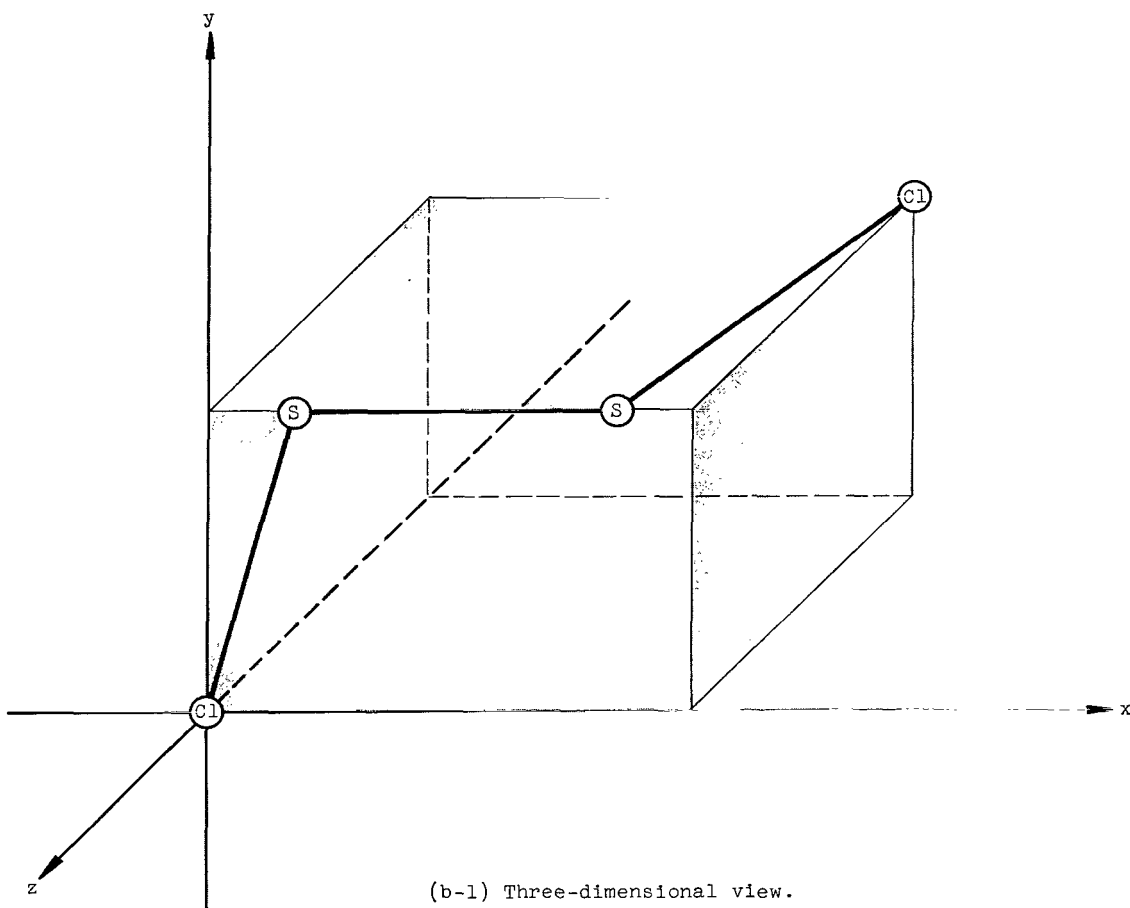


(a-2) Two-dimensional views.

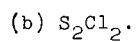
(a) $(\text{NaOH})_2$.

Figure 3. - Diagrams of certain molecules.

CD-7727

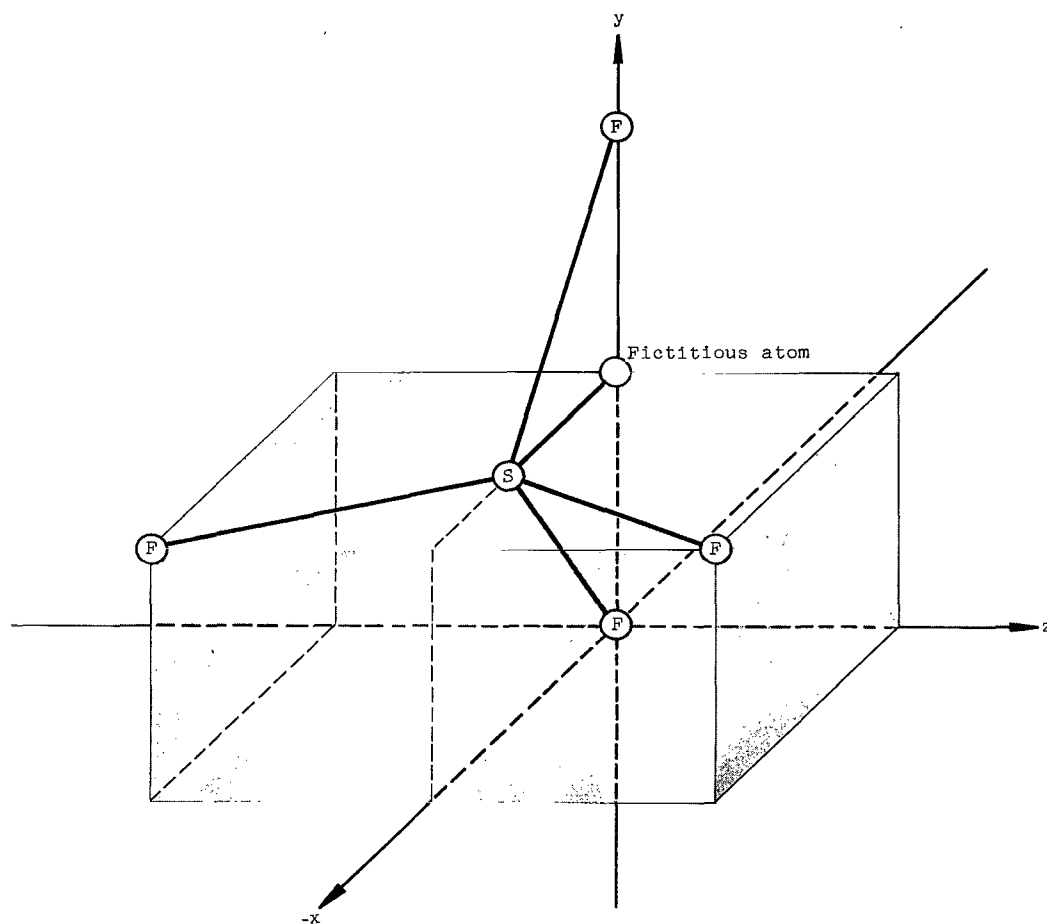


(b-2) Two-dimensional views.

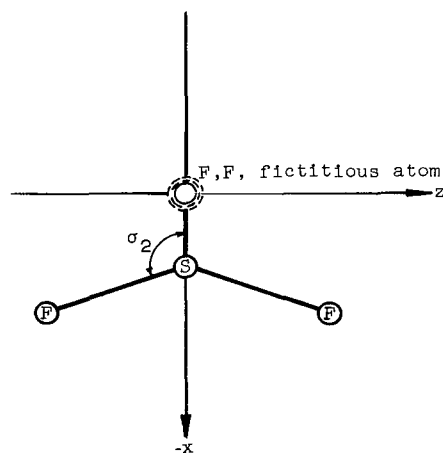
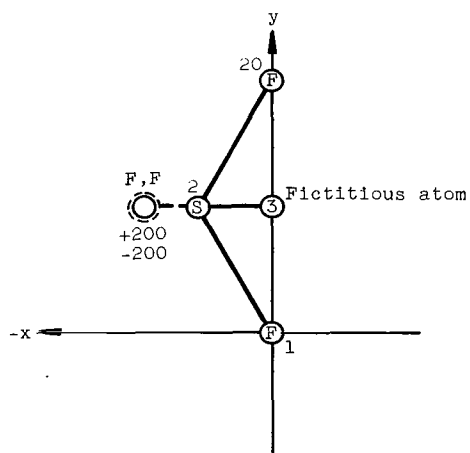


CD-7728

Figure 3. - Continued. Diagrams of certain molecules.



(c-1) Three-dimensional view.



(c-2) Two-dimensional views.

(c) SF_4 .

Figure 3. - Concluded. Diagrams of certain molecules.